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Miller, Charles
Boldog, Ferenc
Li, Li
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Smithson, Glennda
Zerhusen, Bryan
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Tchernev, Velizar
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Rothenberg, Mark

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 <211> 3568
 <212> PRT
 <213> Homo sapiens

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          20          25          30
Leu Phe Pro Glu Thr Ala Pro Gly Ala Pro Gly Ser Ile Pro Ala Pro
          35          40          45
Pro Ala Pro Gly Asp Glu Ala Ala Gly Ser Arg Val Glu Arg Leu Gly
          50          55          60
Gln Ala Phe Arg Val Arg Leu Leu Arg Glu Leu Ser Glu Arg Leu Glu
 65          70          75          80
Leu Val Phe Leu Val Asp Asp Ser Ser Ser Val Gly Glu Val Asn Phe
          85          90          95
Arg Ser Glu Leu Met Phe Val Arg Lys Leu Leu Ser Asp Phe Pro Val
          100          105          110
Val Pro Thr Ala Thr Arg Val Ala Ile Val Thr Phe Ser Ser Lys Asn
          115          120          125
Tyr Val Val Pro Arg Val Asp Tyr Ile Ser Thr Arg Arg Ala Arg Gln
          130          135          140
His Lys Cys Ala Leu Leu Leu Gln Glu Ile Pro Ala Ile Ser Tyr Arg
          145          150          155          160
Gly Gly Gly Thr Tyr Thr Lys Gly Ala Phe Gln Gln Ala Ala Gln Ile
          165          170          175
Leu Leu His Ala Arg Glu Asn Ser Thr Lys Val Val Phe Leu Ile Thr
          180          185          190
Asp Gly Tyr Ser Asn Gly Gly Asp Pro Arg Pro Ile Ala Ala Ser Leu
          195          200          205
Arg Asp Ser Gly Val Glu Ile Phe Thr Phe Gly Ile Trp Gln Gly Asn
          210          215          220
Ile Arg Glu Leu Asn Asp Met Ala Ser Thr Pro Lys Glu Glu His Cys
          225          230          235          240
Tyr Leu Leu His Ser Phe Glu Glu Phe Glu Ala Leu Ala Arg Arg Ala
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His	Cys	Ser	Tyr	Leu	Cys	Asp	Glu	Gly	Lys	Asp	Cys	Cys	Asp	Arg	Met
		275					280					285			
Gly	Ser	Cys	Lys	Cys	Gly	Thr	His	Thr	Gly	His	Phe	Glu	Cys	Ile	Cys
	290					295					300				
Glu	Lys	Gly	Tyr	Tyr	Gly	Lys	Gly	Leu	Gln	Tyr	Glu	Cys	Thr	Ala	Cys
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Pro	Ser	Gly	Thr	Tyr	Lys	Pro	Glu	Ala	Ser	Pro	Gly	Gly	Ile	Ser	Ser
				325					330					335	
Cys	Ile	Pro	Cys	Pro	Asp	Glu	Asn	His	Thr	Ser	Pro	Pro	Gly	Ser	Thr
			340					345					350		
Ser	Pro	Glu	Asp	Cys	Val	Cys	Arg	Glu	Gly	Tyr	Arg	Ala	Ser	Gly	Gln
		355					360					365			
Thr	Cys	Glu	Leu	Val	His	Cys	Pro	Ala	Leu	Lys	Pro	Pro	Glu	Asn	Gly
	370					375					380				
Tyr	Phe	Ile	Gln	Asn	Thr	Cys	Asn	Asn	His	Phe	Asn	Ala	Ala	Cys	Gly
385					390					395					400
Val	Arg	Cys	His	Pro	Gly	Phe	Asp	Leu	Val	Gly	Ser	Ser	Ile	Ile	Leu
				405					410					415	
Cys	Leu	Pro	Asn	Gly	Leu	Trp	Ser	Gly	Ser	Glu	Ser	Tyr	Cys	Arg	Val
			420					425					430		
Arg	Thr	Cys	Pro	His	Leu	Arg	Gln	Pro	Lys	His	Gly	His	Ile	Ser	Cys
		435					440					445			
Ser	Thr	Arg	Glu	Met	Leu	Tyr	Lys	Thr	Thr	Cys	Leu	Val	Ala	Cys	Asp
	450					455					460				
Glu	Gly	Tyr	Arg	Leu	Glu	Gly	Ser	Asp	Lys	Leu	Thr	Cys	Gln	Gly	Asn
465					470					475					480
Ser	Gln	Trp	Asp	Gly	Pro	Glu	Pro	Arg	Cys	Val	Glu	Arg	His	Cys	Ser
				485					490					495	
Thr	Phe	Gln	Met	Pro	Lys	Asp	Val	Ile	Ile	Ser	Pro	His	Asn	Cys	Gly
			500					505					510		
Lys	Gln	Pro	Ala	Lys	Phe	Gly	Thr	Ile	Cys	Tyr	Val	Ser	Cys	Arg	Gln
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Gly	Phe	Ile	Leu	Ser	Gly	Val	Lys	Glu	Met	Leu	Arg	Cys	Thr	Thr	Ser
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Gly	Lys	Trp	Asn	Val	Gly	Val	Gln	Ala	Ala	Val	Cys	Lys	Asp	Val	Glu
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Ala	Pro	Gln	Ile	Asn	Cys	Pro	Lys	Asp	Ile	Glu	Ala	Lys	Thr	Leu	Glu
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Gln	Gln	Asp	Ser	Ala	Asn	Val	Thr	Trp	Gln	Ile	Pro	Thr	Ala	Lys	Asp
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Pro	Tyr	Leu	Phe	Pro	Ile	Gly	Asp	Val	Ala	Ile	Val	Tyr	Thr	Ala	Thr
	610					615					620				
Asp	Leu	Ser	Gly	Asn	Gln	Ala	Ser	Cys	Ile	Phe	His	Ile	Lys	Val	Ile
625					630					635					640
Asp	Ala	Glu	Pro	Pro	Val	Ile	Asp	Trp	Cys	Arg	Ser	Pro	Pro	Pro	Val
				645					650					655	
Gln	Val	Ser	Glu	Lys	Val	His	Ala	Ala	Ser	Trp	Asp	Glu	Pro	Gln	Phe
			660					665					670		
Ser	Asp	Asn	Ser	Gly	Ala	Glu	Leu	Val	Ile	Thr	Arg	Ser	His	Thr	Gln
		675					680					685			
Gly	Asp	Leu	Phe	Pro	Gln	Gly	Glu	Thr	Ile	Val	Gln	Tyr	Thr	Ala	Thr
	690					695					700				
Asp	Pro	Ser	Gly	Asn	Asn	Arg	Ile	Cys	Asp	Ile	His	Ile	Val	Met	Lys
705					710					715					720
Gly	Ser	Pro	Cys	Glu	Ile	Pro	Phe	Thr	Pro	Val	Asn	Gly	Asp	Phe	Ile
				725					730					735	
Cys	Thr	Pro	Asp	Asn	Thr	Gly	Val	Asn	Cys	Thr	Leu	Thr	Cys	Leu	Glu
			740					745					750		
Gly	Tyr	Asp	Phe	Thr	Glu	Gly	Ser	Thr	Asp	Lys	Tyr	Tyr	Cys	Ala	Tyr
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Ala	Lys	Lys	Arg	Phe	Ala	Asn	His	Gly	Phe	Lys	Ser	Phe	Glu	Met	Phe
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Tyr	Lys	Ala	Ala	Arg	Cys	Asp	Asp	Ser	Asp	Leu	Met	Lys	Lys	Phe	Ser
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Glu	Ala	Phe	Glu	Thr	Thr	Leu	Gly	Lys	Met	Val	Pro	Ser	Phe	Cys	Ser
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Asp	Ala	Glu	Asp	Ile	Asp	Cys	Arg	Leu	Glu	Glu	Asn	Leu	Thr	Lys	Lys
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Tyr	Cys	Leu	Glu	Tyr	Asn	Tyr	Asp	Tyr	Glu	Asn	Gly	Phe	Ala	Ile	Gly
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Phe	Leu	Asp	Thr	Val	Gln	Glu	Thr	Ala	Thr	Ser	Ile	Gly	Asn	Ala	Lys
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Leu	Ile	Phe	Asn	Ile	Thr	Ala	Ser	Val	Pro	Leu	Pro	Asp	Glu	Arg	Asn
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 Ser Phe Gln Leu Ala Ser Glu Ile Leu Ile Ala Asp Ser Asn Ser Leu
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 Glu Thr Lys Lys Ala Ser Pro Phe Cys Arg Pro Gly Ser Val Leu Arg
 980 985 990
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 1330 1335 1340
 Glu Cys Leu Ser Gln Pro Cys Lys Asn Gly Ala Thr Cys Lys Asp Gly
 1345 1350 1355 1360
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 Tyr Val Asn Gly Arg Glu Lys Ile Thr Asn Cys Pro Ser Val Asn Asp
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 Pro Val Gln Tyr Cys Leu Asn Gln Gly Gln Trp Thr Gln Pro Leu Pro
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 His Cys Glu Arg Ile Arg Cys Gly Val Pro Pro Pro Leu Glu Asn Gly
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 1925 1930 1935

Pro Ser Ile Ile Glu Cys Thr Ala Ser Gly Ile Trp Asp Arg Ala Pro
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 Pro Ala Cys His Leu Val Phe Cys Gly Glu Pro Pro Ala Ile Lys Asp
 1955 1960 1965
 Ala Val Ile Thr Gly Asn Asn Phe Thr Phe Arg Asn Thr Val Thr Tyr
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 Phe Val Leu Asn Thr Ser Ala Lys Ile Glu Cys Met Arg Gly Gly Gln
 2115 2120 2125
 Trp Asn Pro Ser Pro Met Ser Ile Gln Cys Ile Pro Val Arg Cys Gly
 2130 2135 2140
 Glu Pro Pro Ser Ile Met Asn Gly Tyr Ala Ser Gly Ser Asn Tyr Ser
 2145 2150 2155 2160
 Phe Gly Ala Met Val Ala Tyr Ser Cys Asn Lys Gly Phe Tyr Ile Lys
 2165 2170 2175
 Gly Glu Lys Lys Ser Thr Cys Glu Ala Thr Gly Gln Trp Ser Ser Pro
 2180 2185 2190
 Ile Pro Thr Cys His Pro Val Ser Cys Gly Glu Pro Pro Lys Val Glu
 2195 2200 2205
 Asn Gly Phe Leu Glu His Thr Thr Gly Arg Ile Phe Glu Ser Glu Val
 2210 2215 2220
 Arg Tyr Gln Cys Asn Pro Gly Tyr Lys Ser Val Gly Ser Pro Val Phe
 2225 2230 2235 2240
 Val Cys Gln Ala Asn Arg His Trp His Ser Glu Ser Pro Leu Met Cys
 2245 2250 2255
 Val Pro Leu Asp Cys Gly Lys Pro Pro Pro Ile Gln Asn Gly Phe Met
 2260 2265 2270

Lys Gly Glu Asn Phe Glu Val Gly Ser Lys Val Gln Phe Phe Cys Asn
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 Gly Lys Trp Asn Lys Lys Ser Asn Pro Lys Cys Met Pro Ala Lys Cys
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 Gly Tyr Val Leu Asn Gly Thr Glu Arg Arg Thr Cys Gln Asp Asp Lys
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 3075 3080 3085
 Ser Gly Tyr Val Ile Gln Gly Ser Ser Asp Leu Ile Cys Thr Glu Lys
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 Glu Arg Ile Ser Cys Ser Pro Lys Lys Cys Pro Leu Pro Glu Asn Ile
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 Thr His Ile Leu Val His Gly Asp Asp Phe Ser Val Asn Arg Gln Val
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 Ser Gly Tyr Met Leu Glu Gly Phe Leu Arg Ser Val Cys Leu Glu Asn
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 3555 3560 3565

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 <211> 870
 <212> DNA
 <213> Homo sapiens

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<400> 9

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<211> 209

<212> PRT

<213> Homo sapiens

<400> 10

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Tyr Gln Trp Ala Pro Ile Leu Ala Asn Phe Val His Ile Ile Ile Val
      35          40          45

Ile Leu Gly Leu Phe Gly Thr Ile Gln Tyr Arg Leu Arg Tyr Val Met
      50          55          60

Val Tyr Thr Leu Trp Ala Ala Val Trp Val Thr Trp Asn Val Phe Ile
      65          70          75          80

Ile Cys Phe Tyr Leu Glu Val Gly Gly Leu Leu Lys Asp Ser Glu Leu
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Leu Thr Phe Ser Leu Ser Arg His Arg Ser Trp Trp Arg Glu Arg Trp
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Pro Gly Cys Leu His Glu Glu Val Pro Ala Val Gly Leu Gly Ala Pro
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His Gly Gln Ala Leu Val Ser Gly Ala Gly Cys Ala Leu Glu Pro Ser
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Tyr Val Glu Ala Leu His Ser Cys Leu Gln Ile Leu Ile Ala Leu Leu
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Gly Phe Val Cys Gly Cys Gln Val Val Ser Val Phe Thr Glu Glu Glu
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Asp Ser Phe Asp Phe Ile Gly Gly Phe Asp Pro Phe Pro Leu Tyr His
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Val Asn Glu Lys Pro Ser Ser Leu Leu Ser Lys Gln Val Tyr Leu Pro
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 <212> DNA
 <213> Homo sapiens

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 <211> 197
 <212> PRT
 <213> Homo sapiens

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 35 40 45
 Arg Tyr Val Met Val Tyr Thr Leu Trp Ala Ala Val Trp Val Thr Trp
 50 55 60
 Asn Val Phe Ile Ile Cys Phe Tyr Leu Glu Val Gly Gly Leu Leu Lys
 65 70 75 80
 Asp Ser Glu Leu Leu Thr Phe Ser Leu Ser Arg His Arg Ser Trp Trp
 85 90 95
 Arg Glu Arg Trp Pro Gly Cys Leu His Glu Glu Val Pro Ala Val Gly
 100 105 110
 Leu Gly Ala Pro His Gly Gln Ala Leu Val Ser Gly Ala Gly Cys Ala
 115 120 125
 Leu Glu Pro Ser Tyr Val Glu Ala Leu His Ser Cys Leu Gln Ile Leu
 130 135 140
 Ile Ala Leu Leu Gly Phe Val Cys Gly Cys Gln Val Val Ser Val Phe
 145 150 155 160
 Thr Glu Glu Glu Asp Ser Phe Asp Phe Ile Gly Gly Phe Asp Pro Phe
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 Pro Leu Tyr His Val Asn Glu Lys Pro Ser Ser Leu Leu Ser Lys Gln

180

185

190

Val Tyr Leu Pro Ala
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<211> 909
<212> DNA
<213> Homo sapiens

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<212> PRT
<213> Homo sapiens

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35 40 45
Arg Ile Leu Leu Gly Tyr Asp Gln Gln Ser His Pro Thr Glu His Ser
50 55 60
Lys Gln Met Thr Val Asn Lys Ile Met Val His Ala Asp Tyr Asn Glu
65 70 75 80
Leu His Arg Met Gly Ser Asp Ile Thr Leu Leu Gln Leu His Arg His
85 90 95
Val Glu Phe Ser Ser His Ile Leu Pro Ala Cys Leu Pro Glu Pro Thr
100 105 110
Thr Trp Leu Ala Pro Asp Ser Ser Cys Trp Ile Ser Gly Trp Gly Met
115 120 125
Val Thr Glu Asp Val Phe Leu Pro Glu Pro Phe Gln Leu Gln Glu Ala
130 135 140

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Glu Val Gly Val Met Asp Asn Thr Val Cys Gly Ser Phe Phe Gln Pro
 145 150 155 160
 Gln Tyr Pro Gly Gln Pro Ser Ser Ser Asp Tyr Thr Ile His Glu Asp
 165 170 175
 Met Leu Cys Ala Gly Asp Leu Ile Thr Gly Lys Ala Ile Cys Arg Arg
 180 185 190
 Asp Ser Arg Gly Pro Leu Val Cys Pro Leu Asn Gly Thr Trp Phe Leu
 195 200 205
 Met Gly Leu Ser Ser Trp Ser Leu Asp Cys Cys Ser Pro Val Gly Pro
 210 215 220
 Arg Val Phe Thr Arg Leu Pro Tyr Phe Thr Asn Trp Ile Ser Gln Lys
 225 230 235 240
 Lys Arg Glu Ser Thr Pro Pro Asp Pro Ala Leu Ala Pro Pro Gln Glu
 245 250 255
 Thr Pro Pro Ala Leu Asp Ser Met Thr Ser Gln Gly Ile Val His Lys
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 275 280 285
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 <212> DNA
 <213> Homo sapiens

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<210> 16
 <211> 349
 <212> PRT
 <213> Homo sapiens

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 35 40 45
 Arg Ala Ile Cys Gln Ser Arg Pro Asp Ala Ile Ile Val Ile Gly Glu
 50 55 60
 Gly Ser Gln Met Gly Leu Asp Glu Cys Gln Phe Gln Phe Arg Asn Gly
 65 70 75 80
 Arg Trp Asn Cys Ser Ala Leu Gly Glu Arg Thr Val Phe Gly Lys Glu
 85 90 95
 Leu Lys Val Gly Ser Arg Asp Gly Ala Phe Thr Tyr Ala Ile Ile Ala
 100 105 110
 Ala Gly Val Ala His Ala Ile Thr Ala Ala Cys Thr His Gly Asn Leu
 115 120 125
 Ser Asp Cys Gly Cys Asp Lys Glu Lys Gln Gly Gln Tyr His Arg Asp
 130 135 140
 Glu Gly Trp Lys Trp Gly Gly Cys Ser Ala Asp Ile Arg Tyr Gly Ile
 145 150 155 160
 Gly Phe Ala Lys Val Phe Val Asp Ala Arg Glu Ile Met Lys Asn Ala
 165 170 175
 Arg Arg Leu Met Asn Leu His Asn Asn Glu Ala Gly Arg Lys Val Leu
 180 185 190
 Glu Asp Arg Met Gln Leu Glu Cys Lys Cys His Gly Val Ser Gly Ser
 195 200 205
 Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Lys Phe Arg Glu Val
 210 215 220
 Gly His Leu Leu Lys Glu Lys Tyr Asn Ala Ala Val Gln Val Glu Val
 225 230 235 240
 Val Arg Ala Ser Arg Leu Arg Gln Pro Thr Phe Leu Arg Ile Lys Gln
 245 250 255
 Leu Arg Ser Tyr Arg Lys Pro Met Lys Thr Asp Leu Val Tyr Ile Glu
 260 265 270
 Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly Ser Val Gly
 275 280 285

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Thr Gln Gly Arg Ala Cys Asn Lys Thr Ala Pro Gln Ala Ser Gly Cys
 290 295 300
 Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln Tyr Ala Arg
 305 310 315 320
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 325 330 335
 Asn Thr Cys Ser Glu Arg Thr Glu Met Tyr Thr Cys Lys
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<210> 17
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 <212> DNA
 <213> Homo sapiens

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<210> 18
 <211> 322
 <212> PRT
 <213> Homo sapiens

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 35 40 45
 Arg Asn Gly Arg Trp Asn Cys Ser Ala Leu Gly Glu Arg Thr Val Phe
 50 55 60
 Gly Lys Glu Leu Lys Val Gly Ser Arg Glu Ala Ala Phe Thr Tyr Ala
 65 70 75 80
 Ile Ile Ala Ala Gly Val Ala His Ala Ile Thr Ala Ala Cys Thr Gln
 85 90 95

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Gly Asn Leu Ser Asp Cys Gly Cys Asp Lys Glu Lys Gln Gly Gln Tyr
100 105 110

His Arg Asp Glu Gly Trp Lys Trp Gly Gly Cys Ser Ala Asp Ile Arg
115 120 125

Tyr Gly Ile Gly Phe Ala Lys Val Phe Val Asp Ala Arg Glu Ile Lys
130 135 140

Gln Asn Ala Arg Thr Leu Met Asn Leu His Asn Asn Glu Ala Gly Arg
145 150 155 160

Lys Ile Leu Glu Glu Asn Met Lys Leu Glu Cys Lys Cys His Gly Val
165 170 175

Ser Gly Ser Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Gln Phe
180 185 190

Arg Glu Leu Gly Tyr Val Leu Lys Asp Lys Tyr Asn Glu Ala Val His
195 200 205

Val Glu Pro Val Arg Ala Ser Arg Asn Lys Arg Pro Thr Phe Leu Lys
210 215 220

Ile Lys Lys Pro Leu Ser Tyr Arg Lys Pro Met Asp Thr Asp Leu Val
225 230 235 240

Tyr Ile Glu Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly
245 250 255

Ser Val Gly Thr Gln Gly Arg Ala Cys Asn Lys Thr Ala Pro Gln Ala
260 265 270

Ser Gly Cys Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln
275 280 285

Tyr Ala Arg Val Trp Gln Cys Asn Cys Lys Phe His Trp Cys Cys Tyr
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Val Lys Cys Asn Thr Cys Ser Glu Arg Thr Glu Met Tyr Thr Cys Lys
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Leu Glu

<210> 19
<211> 3731
<212> DNA
<213> Homo sapiens

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 Glu Ala Gln Cys Gly Tyr His Gly Ala Ser Pro Thr Leu Gly Ala Pro
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 Phe Ala Cys Asp Phe Glu Gln Asp Pro Cys Gly Trp Arg Asp Ile Ser
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 Thr Ser Gly Tyr Ser Trp Leu Arg Asp Arg Ala Gly Ala Ala Leu Glu
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 Trp Gly Pro Gly Trp Gln Glu Leu Ala Val Thr Thr Gly Arg Ile Arg
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 Thr Pro Gln Ala Asn Cys Pro Pro Gly His His His Cys Gln Asn Lys
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 Val Cys Val Glu Pro Gln Gln Leu Cys Asp Gly Glu Asp Asn Cys Gly
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Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys
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Glu 465	Gln	Leu	Cys	Ile	Asn 470	Tyr	Thr	Asn	Glu	Lys 475	Leu	Gln	Gln	Leu	Phe 480
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Asp	Glu 530	Glu	Cys	Trp	Phe	Pro 535	Lys	Ala	Thr	Asp	Lys 540	Ser	Phe	Val	Glu
Lys 545	Val	Met	Gln	Glu	Gln 550	Gly	Thr	His	Pro	Lys 555	Phe	Gln	Lys	Pro	Lys 560
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Tyr	Glu	Ile	Leu	Thr 725	Pro	Asn	Ser	Ile	Pro 730	Lys	Gly	Phe	Met	Asp 735	Gly
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1395 1400 1405
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Leu Val Asp Leu Asp His Gln Arg Gln Ser Ala Cys Asn Leu Glu Lys
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 Ser Ile Arg Gly Thr Leu Phe Val Pro Gln Asn Ser Gly Leu Gly Glu
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 Ser Met Phe Phe Tyr Asn Asp Leu Val Asn Gly Thr Thr Leu Gln Thr
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Arg Leu Gly Ser Lys Leu Leu Ile Thr Ala Ser Gln Asp Pro Leu Gln
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Pro Thr Glu Thr Arg Phe Val Asp Gly Arg Ala Ile Leu Gln Trp Asp
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Ala Lys 50 Ile Ile Ser Ser Asp 55 Ile Ile Ser Thr Asn 60 Gly Ile Val His
Ile 65 Ile Asp Lys Leu Leu 70 Ser Pro Lys Asn Leu 75 Leu Ile Thr Pro Lys 80
Asp Asn Ser Gly Arg 85 Ile Leu Gln Asn Leu 90 Thr Thr Leu Ala Thr 95 Asn
Asn Gly Tyr Ile 100 Lys Phe Ser Asn Leu 105 Ile Gln Asp Ser Gly 110 Leu Leu
Ser Val Ile 115 Thr Asp Pro Ile His 120 Thr Pro Val Thr Leu 125 Phe Trp Pro
Thr Asp Gln Ala Leu His Ala Leu Pro Ala Glu Gln Gln Asp Phe Leu
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Val	Ile	Arg	Asp	Ala	Lys	Val	Leu	Ala	Val	Asp	Leu	Pro	Thr	Ser	Thr		
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Ala	Trp	Lys	Thr	Leu	Gln	Gly	Ser	Glu	Leu	Ser	Val	Lys	Cys	Gly	Ala		
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Gly	Arg	Asp	Ile	Gly	Asp	Leu	Phe	Leu	Asn	Gly	Gln	Thr	Cys	Arg	Ile		
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Val	Gln	Arg	Glu	Leu	Leu	Phe	Asp	Leu	Gly	Val	Ala	Tyr	Gly	Ile	Asp		
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Cys	Leu	Leu	Ile	Asp	Pro	Thr	Leu	Gly	Gly	Arg	Cys	Asp	Thr	Phe	Thr		
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Thr	Phe	Asp	Ala	Ser	Gly	Glu	Cys	Gly	Ser	Cys	Val	Asn	Thr	Pro	Ser		
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Cys	Ser	Ala	His	Ala	Thr	Cys	Lys	Glu	Asn	Asn	Thr	Cys	Glu	Cys	Asn		
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Val	Phe	His	Leu	Arg	Ser	Pro
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Gly	Trp	Leu	Glu	Thr	Gly	Arg
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Gln	Asn	Cys	Gly	Ser	Gly	Val
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Pro	Asn	Lys	Ser	Glu	Met	Trp
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Ser	Gly	Asn	Leu	Leu	Gln	Val
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Phe	Leu	Thr	Glu	Val	Leu	Ala
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Ala	Phe	Leu	Glu	His	Leu	Thr
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Val	Pro	Gln	Asn	Ser	Gly	Leu
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Asp	Ile	Glu	His	His	Leu	Ala
				725		
Leu	Val	Asn	Gly	Thr	Thr	Leu
				740		
Ile	Thr	Ala	Ser	Gln	Asp	Pro
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Asp	Gly	Arg	Ala	Ile	Leu	Gln
				770		
Ile	His	Val	Ile	Ser	Arg	Pro
				785		
Leu	Thr	His	Thr	Gly	Leu	Gly

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Lys	Glu	Cys	Val	Tyr	Ile	His	Asp	Pro	Thr	Gly	Leu	Asn	Val	Leu	Lys
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 50 55 60
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 100 105 110
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 115 120 125
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 130 135 140
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 145 150 155 160
 Val Phe His Val Val Thr Gly Leu Arg Trp Gln Ala Pro Ser Gly Thr
 165 170 175
 Pro Gly Asp Pro Lys Arg Thr Ile Gly Gln Ile Leu Ala Ser Thr Glu
 180 185 190
 Ala Phe Ser Arg Phe Glu Thr Ile Leu Glu Asn Cys Gly Leu Pro Ser
 195 200 205
 Ile Leu Asp Gly Pro Gly Pro Phe Thr Val Phe Ala Pro Ser Asn Glu
 210 215 220
 Ala Val Asp Ser Leu Arg Asp Gly Arg Leu Ile Tyr Leu Phe Thr Ala
 225 230 235 240
 Gly Leu Ser Lys Leu Gln Glu Leu Val Arg Tyr His Ile Tyr Asn His
 245 250 255
 Gly Gln Leu Thr Val Glu Lys Leu Ile Ser Lys Gly Arg Ile Leu Thr
 260 265 270

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Met Ala Asn Gln Val Leu Ala Val Asn Ile Ser Glu Glu Gly Arg Ile
 275 280 285
 Leu Leu Gly Pro Glu Gly Val Pro Leu Gln Arg Val Asp Val Met Ala
 290 295 300
 Ala Asn Gly Val Ile His Met Leu Asp Gly Ile Leu Leu Pro Pro Thr
 305 310 315 320
 Ile Leu Pro Ile Leu Pro Lys His Cys Ser Glu Glu Gln His Lys Ile
 325 330 335
 Val Ala Gly Ser Cys Val Asp Cys Gln Ala Leu Asn Thr Ser Thr Cys
 340 345 350
 Pro Pro Asn Ser Val Lys Leu Asp Ile Phe Pro Lys Glu Cys Val Tyr
 355 360 365
 Ile His Asp Pro Thr Gly Leu Asn Val Leu Lys Lys Gly Cys Ala Ser
 370 375 380
 Tyr Cys Asn Gln Thr Ile Met Glu Gln Gly Cys Cys Lys Gly Phe Phe
 385 390 395 400
 Gly Pro Asp Cys Thr Gln Cys Pro Gly Gly Phe Ser Asn Pro Cys Tyr
 405 410 415
 Gly Lys Gly Asn Cys Ser Asp Gly Ile Gln Gly Asn Gly Ala Cys Leu
 420 425 430
 Cys Phe Pro Asp Tyr Lys Gly Ile Ala Cys His Ile Cys Ser Asn Pro
 435 440 445
 Asn Lys His Gly Glu Gln Cys Gln Glu Asp Cys Gly Cys Val His Gly
 450 455 460
 Leu Cys Asp Asn Arg Pro Gly Ser Gly Gly Val Cys Gln Gln Gly Thr
 465 470 475 480
 Cys Ala Pro Gly Phe Ser Gly Arg Phe Cys Asn Glu Ser Met Gly Asp
 485 490 495
 Cys Gly Pro Thr Gly Leu Ala Gln His Cys His Leu His Ala Arg Cys
 500 505 510
 Val Ser Gln Glu Gly Val Ala Arg Cys Arg Cys Leu Asp Gly Phe Glu
 515 520 525
 Gly Asp Gly Phe Ser Cys Thr Pro Ser Asn Pro Cys Ser His Pro Asp
 530 535 540
 Arg Gly Gly Cys Ser Glu Asn Ala Glu Cys Val Pro Gly Ser Leu Gly
 545 550 555 560
 Thr His His Cys Thr Cys His Lys Gly Trp Ser Gly Asp Gly Arg Val
 565 570 575
 Cys Val Ala Ile Asp Glu Cys Glu Leu Asp Val Arg Gly Gly Cys His
 580 585 590
 Thr Asp Ala Leu Cys Ser Tyr Val Gly Pro Gly Gln Ser Arg Cys Thr
 595 600 605

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Cys Lys Leu Gly Phe Ala Gly Asp Gly Tyr Gln Cys Ser Pro Ile Asp
 610 615 620
 Pro Cys Arg Ala Gly Asn Gly Gly Cys His Gly Leu Glu Leu Glu Ala
 625 630 635 640
 Asn Ala His Phe Ser Ile Phe Tyr Gln Trp Leu Lys Ser Ala Gly Ile
 645 650 655
 Thr Leu Pro Ala Asp Arg Arg Val Thr Ala Leu Val Pro Ser Glu Ala
 660 665 670
 Ala Val Arg Gln Leu Ser Pro Glu Asp Arg Ala Phe Trp Leu Gln Pro
 675 680 685
 Arg Thr Leu Pro Asn Leu Val Arg Ala His Phe Leu Gln Gly Ala Leu
 690 695 700
 Phe Glu Glu Glu Leu Ala Arg Leu Gly Gly Gln Glu Val Ala Thr Leu
 705 710 715 720
 Asn Pro Thr Thr Arg Trp Glu Ile Arg Asn Ile Ser Gly Arg Val Trp
 725 730 735
 Val Gln Asn Ala Ser Val Asp Val Ala Asp Leu Leu Ala Thr Asn Gly
 740 745 750
 Val Leu His Ile Leu Ser Gln Val Leu Leu Pro Pro Arg Gly Asp Val
 755 760 765
 Pro Gly Gly Gln Gly Leu Leu Gln Gln Leu Asp Leu Val Pro Ala Phe
 770 775 780
 Ser Leu Phe Arg Glu Leu Leu Gln His His Gly Leu Val Pro Gln Ile
 785 790 795 800
 Glu Ala Ala Thr Ala Tyr Thr Ile Phe Val Pro Thr Asn Arg Ser Leu
 805 810 815
 Glu Ala Gln Gly Asn Ser Ser His Leu Asp Ala Asp Thr Val Arg His
 820 825 830
 His Val Val Leu Gly Glu Ala Leu Ser Met Glu Thr Leu Arg Lys Gly
 835 840 845
 Gly His Arg Asn Ser Leu Leu Gly Pro Ala His Trp Ile Val Phe Tyr
 850 855 860
 Asn His Ser Gly Gln Pro Glu Val Asn His Val Pro Leu Glu Gly Pro
 865 870 875 880
 Met Leu Glu Ala Pro Gly Arg Ser Leu Ile Gly Leu Ser Gly Val Leu
 885 890 895
 Thr Val Gly Ser Ser Arg Cys Leu His Ser His Ala Glu Ala Leu Arg
 900 905 910
 Glu Lys Cys Val Asn Cys Thr Arg Arg Phe Arg Cys Thr Gln Gly Phe
 915 920 925
 Gln Leu Gln Asp Thr Pro Arg Lys Ser Cys Val Tyr Arg Ser Gly Phe
 930 935 940

Ser Phe Ser Arg Gly Cys Ser Tyr Thr Cys Ala Lys Lys Ile Gln Val
 945 950 955 960
 Pro Asp Cys Cys Pro Gly Phe Phe Gly Thr Leu Cys Glu Pro Cys Pro
 965 970 975
 Gly Gly Leu Gly Gly Val Cys Ser Gly His Gly Gln Cys Gln Asp Arg
 980 985 990
 Phe Leu Gly Ser Gly Glu Cys His Cys His Glu Gly Phe His Gly Thr
 995 1000 1005
 Ala Cys Glu Val Cys Glu Leu Gly Arg Tyr Gly Pro Asn Cys Thr Gly
 1010 1015 1020
 Val Cys Asp Cys Ala His Gly Leu Cys Gln Glu Gly Leu Gln Gly Asp
 1025 1030 1035 1040
 Gly Ser Cys Val Cys Asn Val Gly Trp Gln Gly Leu Arg Cys Asp Gln
 1045 1050 1055
 Lys Ile Thr Ser Pro Gln Cys Pro Arg Lys Cys Asp Pro Asn Ala Asn
 1060 1065 1070
 Cys Val Gln Asp Ser Ala Gly Ala Ser Thr Cys Ala Cys Ala Ala Gly
 1075 1080 1085
 Tyr Ser Gly Asn Gly Ile Phe Cys Ser Glu Val Asp Pro Cys Ala His
 1090 1095 1100
 Gly His Gly Gly Cys Ser Pro His Ala Asn Cys Thr Lys Val Ala Pro
 1105 1110 1115 1120
 Gly Gln Arg Thr Cys Thr Cys Gln Asp Gly Tyr Met Gly Asp Gly Glu
 1125 1130 1135
 Leu Cys Gln Glu Ile Asn Ser Cys Leu Ile His His Gly Gly Cys His
 1140 1145 1150
 Ile His Ala Glu Cys Ile Pro Thr Gly Pro Gln Gln Val Ser Cys Ser
 1155 1160 1165
 Cys Arg Glu Gly Tyr Ser Gly Asp Gly Ile Arg Thr Cys Glu Leu Leu
 1170 1175 1180
 Asp Pro Cys Ser Lys Asn Asn Gly Gly Cys Ser Pro Tyr Ala Thr Cys
 1185 1190 1195 1200
 Lys Ser Thr Gly Asp Gly Gln Arg Thr Cys Thr Cys Asp Thr Ala His
 1205 1210 1215
 Thr Val Gly Asp Gly Leu Thr Cys Arg Ala Arg Val Gly Leu Glu Leu
 1220 1225 1230
 Leu Arg Asp Lys His Ala Ser Phe Phe Ser Leu Arg Leu Leu Glu Tyr
 1235 1240 1245
 Lys Glu Leu Lys Gly Asp Gly Pro Phe Thr Ile Phe Val Pro His Ala
 1250 1255 1260
 Asp Leu Met Ser Asn Leu Ser Gln Asp Glu Leu Ala Arg Ile Arg Ala
 1265 1270 1275 1280

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His Arg Gln Leu Val Phe Arg Tyr His Val Val Gly Cys Arg Arg Leu
 1285 1290 1295
 Arg Ser Glu Asp Leu Leu Glu Gln Gly Tyr Ala Thr Ala Leu Ser Gly
 1300 1305 1310
 His Pro Leu Arg Phe Ser Glu Arg Glu Gly Ser Ile Tyr Leu Asn Asp
 1315 1320 1325
 Phe Ala Arg Val Val Ser Ser Asp His Glu Ala Val Asn Gly Ile Leu
 1330 1335 1340
 His Phe Ile Asp Arg Val Leu Leu Pro Pro Glu Ala Leu His Trp Glu
 1345 1350 1355 1360
 Pro Asp Asp Ala Pro Ile Pro Arg Arg Asn Val Thr Ala Ala Ala Gln
 1365 1370 1375
 Gly Phe Gly Tyr Lys Ile Phe Ser Gly Leu Leu Lys Val Ala Gly Leu
 1380 1385 1390
 Leu Pro Leu Leu Arg Glu Ala Ser His Arg Pro Phe Thr Met Leu Trp
 1395 1400 1405
 Pro Thr Asp Ala Ala Phe Arg Ala Leu Pro Pro Asp Arg Gln Ala Trp
 1410 1415 1420
 Leu Tyr His Glu Asp His Arg Asp Lys Leu Ala Ala Ile Leu Arg Gly
 1425 1430 1435 1440
 His Met Ile Arg Asn Val Glu Ala Leu Ala Ser Asp Leu Pro Asn Leu
 1445 1450 1455
 Gly Pro Leu Arg Thr Met His Gly Thr Pro Ile Ser Phe Ser Cys Ser
 1460 1465 1470
 Arg Thr Arg Pro Gly Glu Leu Met Val Gly Glu Asp Asp Ala Arg Ile
 1475 1480 1485
 Val Gln Arg His Leu Pro Phe Glu Gly Gly Leu Ala Tyr Gly Ile Asp
 1490 1495 1500
 Gln Leu Leu Glu Pro Pro Gly Leu Gly Ala Arg Cys Asp His Phe Glu
 1505 1510 1515 1520
 Thr Arg Pro Leu Arg Leu Asn Thr Cys Ser Ile Cys Gly Leu Glu Pro
 1525 1530 1535
 Pro Cys Pro Glu Gly Ser Gln Glu Gln Gly Ser Pro Glu Ala Cys Trp
 1540 1545 1550
 Arg Phe Tyr Pro Lys Phe Trp Thr Ser Pro Pro Leu His Ser Leu Gly
 1555 1560 1565
 Leu Arg Ser Val Trp Val His Pro Ser Leu Trp Gly Arg Pro Gln Gly
 1570 1575 1580
 Leu Gly Arg Gly Cys His Arg Asn Cys Val Thr Thr Thr Trp Lys Pro
 1585 1590 1595 1600
 Ser Cys Cys Pro Gly His Tyr Gly Ser Glu Cys Gln Ala Cys Pro Gly
 1605 1610 1615

Gly Pro Ser Ser Pro Cys Ser Asp Arg Gly Val Cys Met Asp Gly Met
 1620 1625 1630
 Ser Gly Ser Gly Gln Cys Leu Cys Arg Ser Gly Phe Ala Gly Thr Ala
 1635 1640 1645
 Cys Glu Leu Cys Ala Pro Gly Ala Phe Gly Pro His Cys Gln Ala Cys
 1650 1655 1660
 Arg Cys Thr Val His Gly Arg Cys Asp Glu Gly Leu Gly Gly Ser Gly
 1665 1670 1675 1680
 Ser Cys Phe Cys Asp Glu Gly Trp Thr Gly Pro Arg Cys Glu Val Gln
 1685 1690 1695
 Leu Glu Leu Gln Pro Val Cys Thr Pro Pro Cys Ala Pro Glu Ala Val
 1700 1705 1710
 Cys Arg Ala Gly Asn Ser Cys Glu Cys Ser Leu Gly Tyr Glu Gly Asp
 1715 1720 1725
 Gly Arg Val Cys Thr Val Ala Asp Leu Cys Gln Asp Gly His Gly Gly
 1730 1735 1740
 Cys Ser Glu His Ala Asn Cys Ser Gln Val Gly Thr Met Val Thr Cys
 1745 1750 1755 1760
 Thr Cys Leu Pro Asp Tyr Glu Gly Asp Gly Trp Ser Cys Arg Ala Arg
 1765 1770 1775
 Asn Pro Cys Thr Asp Gly His Arg Gly Gly Cys Ser Glu His Ala Asn
 1780 1785 1790
 Cys Leu Ser Thr Gly Leu Asn Thr Arg Arg Cys Glu Cys His Ala Gly
 1795 1800 1805
 Tyr Val Gly Asp Gly Leu Gln Cys Leu Glu Glu Ser Glu Pro Pro Val
 1810 1815 1820
 Asp Arg Cys Leu Gly Gln Pro Pro Pro Cys His Ser Asp Ala Met Cys
 1825 1830 1835 1840
 Thr Asp Leu His Phe Gln Glu Lys Arg Ala Gly Val Phe His Leu Gln
 1845 1850 1855
 Ala Thr Ser Gly Pro Tyr Gly Leu Asn Phe Ser Glu Ala Glu Ala Ala
 1860 1865 1870
 Cys Glu Ala Gln Gly Ala Val Leu Ala Ser Phe Pro Gln Leu Ser Ala
 1875 1880 1885
 Ala Gln Gln Leu Gly Phe His Leu Cys Leu Met Gly Trp Leu Ala Asn
 1890 1895 1900
 Gly Ser Thr Ala His Pro Val Val Phe Pro Val Ala Asp Cys Gly Asn
 1905 1910 1915 1920
 Gly Arg Val Gly Ile Val Ser Leu Gly Ala Arg Lys Asn Leu Ser Glu
 1925 1930 1935
 Arg Trp Asp Ala Tyr Cys Phe Arg Val Gln Asp Val Ala Cys Arg Cys
 1940 1945 1950

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Arg Asn Gly Phe Val Gly Asp Gly Ile Ser Thr Cys Asn Gly Lys Leu
 1955 1960 1965
 Leu Asp Val Leu Ala Ala Thr Ala Asn Phe Ser Thr Phe Tyr Gly Met
 1970 1975 1980
 Leu Leu Gly Tyr Ala Asn Ala Thr Gln Arg Gly Leu Asp Phe Leu Asp
 1985 1990 1995 2000
 Phe Leu Asp Asp Glu Leu Thr Tyr Lys Thr Leu Phe Val Pro Val Asn
 2005 2010 2015
 Glu Gly Phe Val Asp Asn Met Thr Leu Ser Gly Pro Asp Leu Glu Leu
 2020 2025 2030
 His Ala Ser Asn Ala Thr Leu Leu Ser Ala Asn Ala Ser Gln Gly Lys
 2035 2040 2045
 Leu Leu Pro Ala His Ser Gly Leu Ser Leu Ile Ile Ser Asp Ala Gly
 2050 2055 2060
 Pro Asp Asn Ser Ser Trp Ala Pro Val Ala Pro Gly Thr Val Val Val
 2065 2070 2075 2080
 Ser Arg Ile Ile Val Trp Asp Ile Met Ala Phe Asn Gly Ile Ile His
 2085 2090 2095
 Ala Leu Ala Ser Pro Leu Leu Ala Pro Pro Gln Pro Ala Val Leu Ala
 2100 2105 2110
 Pro Glu Ala Pro Pro Val Ala Ala Gly Val Gly Ala Val Leu Ala Ala
 2115 2120 2125
 Gly Ala Leu Leu Gly Leu Val Ala Gly Ala Leu Tyr Leu Arg Ala Arg
 2130 2135 2140
 Gly Lys Pro Met Gly Phe Gly Phe Ser Ala Phe Gln Ala Glu Asp Asp
 2145 2150 2155 2160
 Ala Asp Asp Asp Phe Ser Pro Trp Gln Glu Gly Thr Asn Pro Thr Leu
 2165 2170 2175
 Val Ser Val Pro Asn Pro Val Phe Gly Ser Asp Thr Phe Cys Glu Pro
 2180 2185 2190
 Phe Asp Asp Ser Leu Leu Glu Glu Asp Phe Pro Asp Thr Gln Arg Ile
 2195 2200 2205
 Leu Thr Val Lys
 2210

<210> 44

<211> 149

<212> PRT

<213> Artificial Sequence

<220>

 <223> Description of Artificial Sequence: Fasciclin
 Domain Sequence

<400> 44

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Ala Gly Thr Val Met Glu Lys Leu Lys Thr Asp Pro Arg Phe Ser Thr
 1 5 10 15
 Leu Val Ala Ala Leu Glu Ala Ala Asp Leu Val Glu Thr Leu Asn Asn
 20 25 30
 Ser Gly Asp Phe Thr Val Phe Ala Pro Thr Asn Asp Ala Phe Gln Lys
 35 40 45
 Leu Pro Ala Gly Asp Leu Lys Thr Leu Asp Glu Leu Leu Asn Lys Glu
 50 55 60
 Asp Ala Lys Gln Leu Ala Lys Ile Leu Thr Tyr His Val Val Ala Gly
 65 70 75 80
 Lys Leu Ser Thr Ala Asp Leu Leu Ser Leu Ser Thr Pro Val Leu Thr
 85 90 95
 Ser Leu Gln Gly Ser Lys Ile Thr Val Ser Gly Lys Asn Asp Thr Glu
 100 105 110
 Leu Leu Lys Asp Val Asn Val Leu Lys Val Asn Asn Ala Thr Val Ile
 115 120 125
 Val Glu Ser Asp Ile Glu Thr Thr Asn Gly Val Ile His Val Ile Asp
 130 135 140
 Arg Val Leu Leu Pro
 145

<210> 45
 <211> 149
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Fasciclin
 domain sequence

<400> 45
 Ala Gly Thr Val Met Glu Lys Leu Lys Thr Asp Pro Arg Phe Ser Thr
 1 5 10 15
 Leu Val Ala Ala Leu Glu Ala Ala Asp Leu Val Glu Thr Leu Asn Asn
 20 25 30
 Ser Gly Asp Phe Thr Val Phe Ala Pro Thr Asn Asp Ala Phe Gln Lys
 35 40 45
 Leu Pro Ala Gly Asp Leu Lys Thr Leu Asp Glu Leu Leu Asn Lys Glu
 50 55 60
 Asp Ala Lys Gln Leu Ala Lys Ile Leu Thr Tyr His Val Val Ala Gly
 65 70 75 80
 Lys Leu Ser Thr Ala Asp Leu Leu Ser Leu Ser Thr Pro Val Leu Thr
 85 90 95
 Ser Leu Gln Gly Ser Lys Ile Thr Val Ser Gly Lys Asn Asp Thr Glu
 100 105 110
 Leu Leu Lys Asp Val Asn Val Leu Lys Val Asn Asn Ala Thr Val Ile
 Page 112

115 120 125
 Val Glu Ser Asp Ile Glu Thr Thr Asn Gly Val Ile His Val Ile Asp
 130 135 140

Arg Val Leu Leu Pro
 145

<210> 46
 <211> 104
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: XLINK domain
 sequence

<400> 46
 Gly Glu Val Phe His Tyr Arg Ala Pro Ser Gly Arg Tyr Lys Leu Thr
 1 5 10 15
 Phe Glu Glu Ala Gln Ala Ala Cys Leu Arg Gln Gly Ala Arg Ile Ala
 20 25 30
 Thr Thr Gly Gln Leu Tyr Ala Ala Trp Lys Gly Gly Phe Asp Gln Cys
 35 40 45
 Asp Ala Gly Trp Leu Ala Asp Gly Ser Val Arg Tyr Pro Ile Val Lys
 50 55 60
 Pro Arg Glu Asn Cys Gly Gly Asp Lys Asp Gly Phe Pro Gly Val Arg
 65 70 75 80
 Thr Tyr Tyr Leu Phe Pro Asn Gln Thr Gly Phe Pro Asp Asp Pro Ser
 85 90 95
 Ser Arg Tyr Asp Val Tyr Cys Phe
 100

<210> 47
 <211> 3567
 <212> PRT
 <213> Mus musculus

<400> 47
 Met Trp Ser Arg Leu Ala Phe Cys Cys Trp Ala Leu Ala Leu Val Ser
 1 5 10 15
 Gly Trp Thr Asn Phe Gln Pro Val Ala Pro Ser Leu Asn Phe Ser Phe
 20 25 30
 Arg Leu Phe Pro Glu Ala Ser Pro Gly Ala Leu Gly Arg Leu Ala Val
 35 40 45
 Pro Pro Ala Ser Ser Glu Glu Glu Ala Ala Gly Ser Lys Val Glu Arg
 50 55 60
 Leu Gly Arg Ala Phe Arg Ser Arg Val Arg Arg Leu Arg Glu Leu Ser
 65 70 75 80
 Gly Ser Leu Glu Leu Val Phe Leu Val Asp Glu Ser Ser Ser Val Gly
 Page 113

85

90

95

Gln Thr Asn Phe Leu Asn Glu Leu Lys Phe Val Arg Lys Leu Leu Ser
 100 105 110
 Asp Phe Pro Val Val Ser Thr Ala Thr Arg Val Ala Ile Val Thr Phe
 115 120 125
 Ser Ser Lys Asn Asn Val Val Ala Arg Val Asp Tyr Ile Ser Thr Ser
 130 135 140
 Arg Ala His Gln His Lys Cys Ala Leu Leu Ser Arg Glu Ile Pro Ala
 145 150 155 160
 Ile Thr Tyr Arg Gly Gly Gly Thr Tyr Thr Lys Gly Ala Phe Gln Gln
 165 170 175
 Ala Ala Gln Ile Leu Arg His Ser Arg Glu Asn Ser Thr Lys Val Ile
 180 185 190
 Phe Leu Ile Thr Asp Gly Tyr Ser Asn Gly Gly Asp Pro Arg Pro Ile
 195 200 205
 Ala Ala Ser Leu Arg Asp Phe Gly Val Glu Ile Phe Thr Phe Gly Ile
 210 215 220
 Trp Gln Gly Asn Ile Arg Glu Leu Asn Asp Met Ala Ser Thr Pro Lys
 225 230 235 240
 Glu Glu His Cys Tyr Leu Leu His Ser Phe Glu Glu Phe Glu Ala Leu
 245 250 255
 Ala Arg Arg Ala Leu His Glu Asp Leu Pro Ser Gly Ser Phe Ile Gln
 260 265 270
 Glu Asp Met Ala Arg Cys Ser Tyr Leu Cys Glu Ala Gly Lys Asp Cys
 275 280 285
 Cys Asp Arg Met Ala Ser Cys Lys Cys Gly Thr His Thr Gly Gln Phe
 290 295 300
 Glu Cys Ile Cys Glu Lys Gly Tyr Tyr Gly Lys Gly Leu Gln His Glu
 305 310 315 320
 Cys Thr Ala Cys Pro Ser Gly Thr Tyr Lys Pro Glu Ala Ser Pro Gly
 325 330 335
 Gly Ile Ser Thr Cys Ile Pro Cys Pro Asp Val Ser His Thr Ser Pro
 340 345 350
 Pro Gly Ser Thr Ser Pro Glu Asp Cys Val Cys Arg Glu Gly Tyr Gln
 355 360 365
 Arg Ser Gly Gln Thr Cys Glu Val Val His Cys Pro Ala Leu Lys Pro
 370 375 380
 Pro Glu Asn Gly Phe Phe Ile Gln Asn Thr Cys Lys Asn His Phe Asn
 385 390 395 400
 Ala Ala Cys Gly Val Arg Cys Arg Pro Gly Phe Asp Leu Val Gly Ser
 405 410 415
 Ser Ile His Leu Cys Gln Pro Asn Gly Leu Trp Ser Gly Thr Glu Ser
 Page 114

420		425		430
Phe Cys Arg Val Arg Thr Cys Pro His Leu Arg Gln Pro Lys His Gly	435	440	445	
His Ile Ser Cys Ser Thr Ala Glu Met Ser Tyr Asn Thr Leu Cys Leu	450	455	460	
Val Thr Cys Asn Glu Gly Tyr Arg Leu Glu Gly Ser Thr Arg Leu Thr	465	470	475	480
Cys Gln Gly Asn Ala Gln Trp Asp Gly Pro Glu Pro Arg Cys Val Glu	485	490	495	
Arg His Cys Ala Thr Phe Gln Lys Pro Lys Gly Val Ile Ile Ser Pro	500	505	510	
Pro Ser Cys Gly Lys Gln Pro Ala Arg Pro Gly Met Thr Cys Gln Leu	515	520	525	
Ser Cys Arg Gln Gly Tyr Ile Leu Ser Gly Val Arg Glu Val Arg Cys	530	535	540	
Ala Thr Ser Gly Lys Trp Ser Ala Lys Val Gln Thr Ala Val Cys Lys	545	550	555	560
Asp Val Glu Ala Pro Gln Ile Ser Cys Pro Asn Asp Ile Glu Ala Lys	565	570	575	
Thr Gly Glu Gln Gln Asp Ser Ala Asn Val Thr Trp Gln Val Pro Thr	580	585	590	
Ala Lys Asp Asn Ser Gly Glu Lys Val Ser Val His Val His Pro Ala	595	600	605	
Phe Thr Pro Pro Tyr Leu Phe Pro Ile Gly Asp Val Ala Ile Thr Tyr	610	615	620	
Thr Ala Thr Asp Ser Ser Gly Asn Gln Ala Ser Cys Thr Phe Tyr Ile	625	630	635	640
Lys Val Ile Asp Val Glu Pro Pro Val Ile Asp Trp Cys Arg Ser Pro	645	650	655	
Pro Pro Ile Gln Val Val Glu Lys Glu His Pro Ala Ser Trp Asp Glu	660	665	670	
Pro Gln Phe Ser Asp Asn Ser Gly Ala Glu Leu Val Ile Thr Ser Ser	675	680	685	
His Thr Gln Gly Asp Met Phe Pro His Gly Glu Thr Val Val Trp Tyr	690	695	700	
Thr Ala Thr Asp Pro Ser Gly Asn Asn Arg Thr Cys Asp Ile His Ile	705	710	715	720
Val Ile Lys Gly Ser Pro Cys Glu Val Pro Phe Thr Pro Val Asn Gly	725	730	735	
Asp Phe Ile Cys Ala Gln Asp Ser Ala Gly Val Asn Cys Ser Leu Ser	740	745	750	
Cys Lys Glu Gly Tyr Asp Phe Thr Glu Gly Ser Thr Glu Lys Tyr Tyr				

1090 1095 1100
 Cys Gly Val Pro Cys Pro Val Gly Glu Phe Ser Arg Ser Gly Leu Thr
 1105 1110 1115 1120
 Pro Cys Tyr Pro Cys Pro Arg Asp Tyr Tyr Gln Pro Asn Ala Gly Lys
 1125 1130 1135
 Ser Phe Cys Leu Ala Cys Pro Phe Tyr Gly Thr Thr Thr Ile Thr Gly
 1140 1145 1150
 Ala Thr Ser Ile Thr Asp Cys Ser Ser Phe Ser Ser Thr Phe Ser Ala
 1155 1160 1165
 Ala Glu Glu Ser Ile Val Pro Leu Val Ala Pro Gly His Ser Gln Asn
 1170 1175 1180
 Lys Tyr Glu Val Ser Ser Gln Val Phe His Glu Cys Phe Leu Asn Pro
 1185 1190 1195 1200
 Cys His Asn Ser Gly Thr Cys Gln Gln Leu Gly Arg Gly Tyr Val Cys
 1205 1210 1215
 Leu Cys Pro Pro Gly Tyr Thr Gly Leu Lys Cys Glu Thr Asp Ile Asp
 1220 1225 1230
 Glu Cys Ser Ser Leu Pro Cys Leu Asn Gly Gly Ile Cys Arg Asp Gln
 1235 1240 1245
 Val Gly Gly Phe Thr Cys Glu Cys Ser Leu Gly Tyr Ser Gly Gln Ile
 1250 1255 1260
 Cys Glu Glu Asn Ile Asn Glu Cys Ile Ser Ser Pro Cys Leu Asn Lys
 1265 1270 1275 1280
 Gly Thr Cys Thr Asp Gly Leu Ala Ser Tyr Arg Cys Thr Cys Val Lys
 1285 1290 1295
 Gly Tyr Met Gly Val His Cys Glu Thr Asp Val Asn Glu Cys Gln Ser
 1300 1305 1310
 Ser Pro Cys Leu Asn Asn Ala Val Cys Lys Asp Gln Val Gly Gly Phe
 1315 1320 1325
 Ser Cys Lys Cys Pro Pro Gly Phe Leu Gly Thr Arg Cys Glu Lys Asn
 1330 1335 1340
 Val Asp Glu Cys Leu Ser Gln Pro Cys Gln Asn Gly Ala Thr Cys Lys
 1345 1350 1355 1360
 Asp Gly Ala Asn Ser Phe Arg Cys Gln Cys Pro Ala Gly Phe Thr Gly
 1365 1370 1375
 Thr His Cys Glu Leu Asn Ile Asn Glu Cys Gln Ser Asn Pro Cys Arg
 1380 1385 1390
 Asn Gln Ala Thr Cys Val Asp Glu Leu Asn Ser Tyr Ser Cys Lys Cys
 1395 1400 1405
 Gln Pro Gly Phe Ser Gly His Arg Cys Glu Thr Glu Gln Pro Ser Gly
 1410 1415 1420
 Phe Asn Leu Asp Phe Glu Val Ser Gly Ile Tyr Gly Tyr Val Leu Leu
 Page 117

1425 1430 1435 1440
 Asp Gly Val Leu Pro Thr Leu His Ala Ile Thr Cys Ala Phe Trp Met
 1445 1450 1455
 Lys Ser Ser Asp Val Ile Asn Tyr Gly Thr Pro Ile Ser Tyr Ala Leu
 1460 1465 1470
 Glu Asp Asp Lys Asp Asn Thr Ser Leu Leu Thr Asp Tyr Asn Gly Trp
 1475 1480 1485
 Val Leu Tyr Val Asn Gly Lys Glu Lys Ile Thr Asn Cys Pro Ser Val
 1490 1495 1500
 Asn Asp Gly Ile Trp His His Ile Ala Ile Thr Trp Thr Ser Thr Gly
 1505 1510 1515 1520
 Gly Ala Trp Arg Val Tyr Ile Asn Gly Glu Leu Ser Asp Gly Gly Thr
 1525 1530 1535
 Gly Leu Ser Ile Gly Lys Ala Ile Pro Gly Gly Gly Ala Leu Val Leu
 1540 1545 1550
 Gly Gln Glu Gln Asp Lys Lys Gly Glu Gly Phe Asn Pro Ala Glu Ser
 1555 1560 1565
 Phe Val Gly Ser Ile Ser Gln Leu Asn Leu Trp Asp Tyr Val Leu Ser
 1570 1575 1580
 Pro Gln Gln Val Lys Leu Leu Ala Ser Ser Cys Pro Glu Glu Leu Ser
 1585 1590 1595 1600
 Arg Gly Asn Val Leu Ala Trp Pro Asp Phe Leu Ser Gly Ile Thr Gly
 1605 1610 1615
 Lys Val Lys Val Asp Ser Ser Ser Met Phe Cys Ser Asp Cys Pro Ser
 1620 1625 1630
 Leu Glu Gly Ser Val Pro His Leu Arg Pro Ala Ser Gly Asn Arg Lys
 1635 1640 1645
 Pro Gly Ser Lys Val Ser Leu Phe Cys Asp Pro Gly Phe Gln Met Val
 1650 1655 1660
 Gly Asn Pro Val Gln Tyr Cys Leu Asn Gln Gly Gln Trp Thr Gln Pro
 1665 1670 1675 1680
 Leu Pro His Cys Glu Arg Ile Arg Cys Gly Leu Pro Pro Ala Leu Glu
 1685 1690 1695
 Asn Gly Phe Tyr Ser Ala Glu Asp Phe His Ala Gly Ser Thr Val Thr
 1700 1705 1710
 Tyr Gln Cys Thr Ser Gly Tyr Tyr Leu Leu Gly Asp Ser Arg Met Phe
 1715 1720 1725
 Cys Thr Asp Asn Gly Ser Trp Asn Gly Ile Ser Pro Ser Cys Leu Asp
 1730 1735 1740
 Val Asp Glu Cys Ala Val Gly Ser Asp Cys Ser Glu His Ala Ser Cys
 1745 1750 1755 1760
 Leu Asn Thr Asn Gly Ser Tyr Val Cys Ser Cys Asn Pro Pro Tyr Thr
 Page 118

1765 1775
 Gly Asp Gly Lys Asn Cys Ala Glu Pro Val Lys Cys Lys Ala Pro Glu
 1780 1785 1790
 Asn Pro Glu Asn Gly His Ser Ser Gly Glu Ile Tyr Thr Val Gly Thr
 1795 1800 1805
 Ala Val Thr Phe Ser Cys Asp Glu Gly His Glu Leu Val Gly Val Ser
 1810 1815 1820
 Thr Ile Thr Cys Leu Glu Thr Gly Glu Trp Asp Arg Leu Arg Pro Ser
 1825 1830 1835 1840
 Cys Glu Ala Ile Ser Cys Gly Val Pro Pro Val Pro Glu Asn Gly Gly
 1845 1850 1855
 Val Asp Gly Ser Ala Phe Thr Tyr Gly Ser Lys Val Val Tyr Arg Cys
 1860 1865 1870
 Asp Lys Gly Tyr Thr Leu Ser Gly Asp Glu Glu Ser Ala Cys Leu Ala
 1875 1880 1885
 Ser Gly Ser Trp Ser His Ser Ser Pro Val Cys Glu Leu Val Lys Cys
 1890 1895 1900
 Ser Gln Pro Glu Asp Ile Asn Asn Gly Lys Tyr Ile Leu Ser Gly Leu
 1905 1910 1915 1920
 Thr Tyr Leu Ser Ile Ala Ser Tyr Ser Cys Glu Asn Gly Tyr Ser Leu
 1925 1930 1935
 Gln Gly Pro Ser Leu Leu Glu Cys Thr Ala Ser Gly Ser Trp Asp Arg
 1940 1945 1950
 Ala Pro Pro Ser Cys Gln Leu Val Ser Cys Gly Glu Pro Pro Ile Val
 1955 1960 1965
 Lys Asp Ala Val Ile Thr Gly Ser Asn Phe Thr Phe Gly Asn Thr Val
 1970 1975 1980
 Ala Tyr Thr Cys Lys Glu Gly Tyr Thr Leu Ala Gly Pro Asp Thr Ile
 1985 1990 1995 2000
 Val Cys Gln Ala Asn Gly Lys Trp Asn Ser Ser Asn His Gln Cys Leu
 2005 2010 2015
 Ala Val Ser Cys Asp Glu Pro Pro Asn Val Asp His Ala Ser Pro Glu
 2020 2025 2030
 Thr Ala His Arg Leu Phe Gly Asp Thr Ala Phe Tyr Tyr Cys Ala Asp
 2035 2040 2045
 Gly Tyr Ser Leu Ala Asp Asn Ser Gln Leu Ile Cys Asn Ala Gln Gly
 2050 2055 2060
 Asn Trp Val Pro Pro Ala Gly Gln Ala Val Pro Arg Cys Ile Ala His
 2065 2070 2075 2080
 Phe Cys Glu Lys Pro Pro Ser Val Ser Tyr Ser Ile Leu Glu Ser Val
 2085 2090 2095
 Ser Lys Ala Lys Phe Ala Ala Gly Ser Val Val Ser Phe Lys Cys Met
 Page 119

2100 2105 2110
 Glu Gly Phe Val Leu Asn Thr Ser Ala Lys Ile Glu Cys Leu Arg Gly
 2115 2120 2125
 Gly Glu Trp Ser Pro Ser Pro Leu Ser Val Gln Cys Ile Pro Val Arg
 2130 2135 2140
 Cys Gly Glu Pro Pro Ser Ile Ala Asn Gly Tyr Pro Ser Gly Thr Asn
 2145 2150 2155 2160
 Tyr Ser Phe Gly Ala Val Val Ala Tyr Ser Cys His Lys Gly Phe Tyr
 2165 2170 2175
 Ile Lys Gly Glu Lys Lys Ser Thr Cys Glu Ala Thr Gly Gln Trp Ser
 2180 2185 2190
 Lys Pro Thr Pro Thr Cys His Pro Val Ser Cys Asn Glu Pro Pro Lys
 2195 2200 2205
 Val Glu Asn Gly Phe Leu Glu His Thr Thr Gly Arg Thr Phe Glu Ser
 2210 2215 2220
 Glu Ala Arg Phe Gln Cys Asn Pro Gly Tyr Lys Ala Ala Gly Ser Pro
 2225 2230 2235 2240
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 Ala Leu Gln Gly Pro Ser Val Leu Lys Cys Leu Pro Ser Gly Gln Trp
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 2385 2390 2395 2400
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 2420 2425 2430
 Glu Cys Val Pro Val Glu Cys Pro Gln Pro Glu Glu Ile Leu Asn Gly
 Page 120

2435 2440 2445
 Ile Ile His Val Gln Gly Leu Ala Tyr Leu Ser Thr Thr Leu Tyr Thr
 2450 2455 2460
 Cys Lys Pro Gly Phe Glu Leu Val Gly Asn Ala Thr Thr Leu Cys Gly
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 Cys Pro Glu Pro Lys Glu Ile Leu Asn Gly Gln Phe Ser Ser Val Ser
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 Phe Gln Tyr Gly Gln Thr Ile Thr Tyr Phe Cys Asp Arg Gly Phe Arg
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 Page 121

2770

2775

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Gly Tyr Ile Leu Asn Gly Ser Lys Lys Arg Thr Cys Gln Glu Asn Arg
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Pro Pro Val Pro Thr Asn Gly Arg Val Lys Gly Glu Glu Tyr Thr Phe
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Gln Lys Glu Ile Thr Tyr Ser Cys Arg Glu Gly Phe Ile Leu Glu Gly
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Ala Arg Ser Arg Ile Cys Leu Thr Asn Gly Ser Trp Ser Gly Ala Thr
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His Cys Leu Glu Gly Tyr Val Leu Gln Gly Ala Pro Arg Leu Thr Cys
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Ser Gly Ser Ser Pro Ser Cys Leu Pro Cys Arg Cys Ser Thr Pro Ile
2995 3000 3005
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Ile Trp Ser Gln Pro Tyr Pro Thr Cys Glu Pro Leu Ser Cys Gly Pro

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Arg Ile Thr Cys Ser Pro Lys Lys Cys Pro Val Pro Ser Asn Met Thr
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Arg Ile Arg Phe His Gly Asp Asp Phe Gln Val Asn Arg Gln Val Ser
3185 3190 3200

Val Ser Cys Ala Glu Gly Phe Thr His Glu Gly Val Asn Trp Ser Thr
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Cys Gln Pro Asp Gly Thr Trp Glu Pro Pro Phe Ser Asp Glu Ser Cys
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Ile Pro Val Val Cys Gly His Pro Glu Ser Pro Ala His Gly Ser Val
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Val Gly Asn Lys His Ser Phe Gly Ser Thr Ile Val Tyr Gln Cys Asp
3250 3255 3260

Pro Gly Tyr Lys Leu Glu Gly Asn Arg Glu Arg Ile Cys Gln Glu Asn
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Thr Pro Ala Glu Phe Pro Asn Gly Lys Ala Val Leu Glu Asn Thr Thr
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Ser Gly Pro Ser Leu Leu Phe Ser Cys His Arg Gly Tyr Thr Leu Glu
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Gly Ser Pro Glu Ala His Cys Thr Ala Asn Gly Thr Trp Asn His Leu
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Thr Pro Leu Cys Lys Pro Asn Pro Cys Pro Val Pro Phe Val Ile Pro
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Ser Ile Lys Cys Arg Glu Gly Phe Leu Leu Lys Gly Asn Gly Val Ile
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Lys Ile Ser Cys Gly Pro Pro Ser His Val Glu Asn Ala Ile Ala Arg
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Gly Val Tyr Tyr Gln Tyr Gly Asp Met Ile Thr Tyr Ser Cys Tyr Ser
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Gly Tyr Met Leu Glu Gly Ser Leu Arg Ser Val Cys Leu Glu Asn Gly

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3445

3455

Thr Trp Thr Pro Ser Pro Val Cys Arg Ala Val Cys Arg Phe Pro Cys
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Gln Asn Gly Gly Val Cys Gln Arg Pro Asn Ala Cys Ser Cys Pro Asp
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Gly Trp Met Gly Arg Leu Cys Glu Glu Pro Ile Cys Ile Leu Pro Cys
3490 3495 3500
Leu Asn Gly Gly Arg Cys Val Ala Pro Tyr Gln Cys Asp Cys Pro Thr
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Gly Trp Thr Gly Ser Arg Cys His Thr Ala Thr Cys Gln Ser Pro Cys
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<400> 48

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Asp Glu Gly Lys Asp Cys Cys Asp Arg Met Gly Ser Cys Lys Cys Gly
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Glu Asn His Thr Ser Pro Pro Gly Ser Thr Ser Pro Glu Asp Cys Val
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Cys Arg Glu Gly Tyr Arg Ala Ser Gly Gln Thr Cys Glu Leu Val His
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Cys Asn Asn His Phe Asn Ala Ala Cys Gly Val Arg Cys His Pro Gly
165 170 175

Phe Asp Leu Val Gly Ser Ser Ile Ile Leu Cys Leu Pro Asn Gly Leu
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 Trp Ser Gly Leu Glu Ser Tyr Cys Arg Val Arg Thr Cys Pro His Leu
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 Arg Gln Pro Lys His Gly His Ile Ser Cys Ser Thr Arg Glu Met Leu
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 Tyr Lys Thr Thr Cys Leu Val Ala Cys Asp Glu Gly Tyr Arg Leu Glu
 225 230 235 240
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 260 265 270
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 Val Lys Glu Met Leu Arg Cys Thr Thr Ser Gly Lys Trp Asn Val Gly
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 370 375 380
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 Ile Asp Trp Cys Arg Ser Pro Pro Val Gln Val Ser Glu Lys Val
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 His Ala Ala Ser Trp Asp Glu Pro Gln Phe Ser Asp Asn Ser Gly Ala
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 Leu Gly Lys Met Val Pro Ser Phe Cys Ser Asp Ala Glu Asp Ile Asp
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 645 650 655
 Glu Thr Ala Thr Ser Ile Gly Asn Ala Lys Ser Ser Arg Ile Lys Arg
 660 665 670
 Ser Ala Pro Leu Ser Asp Tyr Lys Ile Lys Leu Ile Phe Asn Ile Thr
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 Ala Ser Val Pro Leu Pro Asp Glu Arg Asn Asp Thr Leu Glu Trp Glu
 690 695 700
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 Glu Ile Leu Ile Ala Asp Ser Asn Ser Leu Gly Thr Lys Lys Ala Ser
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 770 775 780
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 785 790 795 800
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 Leu Glu Thr Cys Glu Ser Cys Pro Leu Gly Thr Tyr Gln Pro Lys Phe
 835 840 845

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Lys	Phe	Ser	Arg	Ser	Gly	Leu	Met	Pro	Cys	His	Pro	Cys	Pro	Arg	Asp
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Tyr	Tyr	Gln	Pro	Asn	Ala	Gly	Lys	Ala	Phe	Cys	Leu	Ala	Cys	Pro	Phe
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Leu	Lys	Cys	Glu	Thr	Asp	Ile	Asp	Glu	Cys	Ser	Pro	Leu	Pro	Cys	Leu
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Leu	Cys	Ala	Ala	Gly	Phe	Thr	Gly	Ser	His	Cys	Glu	Leu	Asn	Ile	Asn
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Glu	Cys	Gln	Ser	Asn	Pro	Cys	Arg	Asn	Gln	Ala	Thr	Cys	Val	Asp	Glu
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Leu	Asn	Ser	Tyr	Ser	Cys	Lys	Cys	Gln	Pro	Gly	Phe	Ser	Gly	Lys	Arg
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Gly Ile Tyr Gly Tyr Val Met Leu Val Gly Met Leu Pro Ser Leu His
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Gly Thr Pro Ile Ser Tyr Ala Val Asp Asn Gly Ser Asp Asn Thr Leu
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Lys Ile Thr Asn Cys Pro Ser Val Asn Asp Gly Arg Trp His His Ile
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Ser Ser Pro Ser Cys Leu Pro Cys Arg Cys Ser Thr Pro Ile Ile Gln
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Cys Asp Ala Asn Gly Gln Trp Ser Asp Val Pro Leu Cys Glu His Ala
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 Asn Lys His Ser Phe Gly Ser Thr Ile Val Tyr Gln Cys Asp Pro Gly
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 Trp Ser Gly Glu Val Ala Val Cys Arg Glu Asn Arg Cys Glu Thr Pro
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 Pro Ser Leu Leu Phe Ser Cys His Arg Gly Tyr Thr Leu Glu Gly Ser
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 Pro Glu Ala His Cys Thr Ala Asn Gly Thr Trp Asn His Leu Thr Pro
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 465 470 475 480
 Lys Cys Arg Glu Gly Phe Leu Leu Lys Gly Asn Gly Val Ile Thr Cys
 485 490 495

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Ser Pro Asp Glu Thr Trp Thr His Thr Asn Ala Arg Cys Glu Lys Ile
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 Leu Pro Thr Ile Pro Asn Ala Ile Val Leu Glu Gly Ser Leu Ser Glu
 100 105 110
 Asp Asn Val Val Thr Tyr Ser Cys Arg Pro Gly Tyr Thr Met Gln Gly
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Asn	Gly	Lys	Ala	Val	Leu	Glu	Asn	Thr	Thr	Ser	Gly	Pro	Ser	Leu	Leu
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Phe	Ser	Cys	His	Arg	Gly	Tyr	Thr	Leu	Glu	Gly	Ser	Pro	Glu	Ala	His
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Cys	Thr	Ala	Asn	Gly	Thr	Trp	Asn	His	Leu	Thr	Pro	Leu	Cys	Lys	Pro
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Pro	Ser	His	Val	Glu	Asn	Ala	Ile	Ala	Arg	Gly	Val	Tyr	Tyr	Gln	Tyr

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 Gly Asp Met Ile Thr Tyr Ser Cys Tyr Ser Gly Tyr Met Leu Glu Gly
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 Val Cys Arg Ala Val Cys Arg Phe Pro Cys Gln Asn Gly Gly Val Cys
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 Gln Arg Pro Asn Ala Cys Ser Cys Pro Asp Gly Trp Met Gly Arg Leu
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 Val Ala Pro Tyr Gln Cys Asp Cys Pro Thr Gly Trp Thr Gly Ser Arg
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 Cys His Thr Ala Thr Cys Gln Ser Pro Cys Leu Asn Gly Gly Lys Cys
 565 570 575
 Ile Arg Pro Asn Arg Cys His Cys Leu Ser Ala Trp Thr Gly His Asp
 580 585 590
 Cys Ser Arg Lys Arg Arg Ala Gly Leu
 595 600

<210> 51
 <211> 481
 <212> PRT
 <213> Homo sapiens

<400> 51
 Met Lys Gly Glu Asn Phe Glu Val Gly Ser Lys Val Gln Phe Phe Cys
 1 5 10 15
 Asn Glu Gly Tyr Glu Leu Val Gly Asp Ser Ser Trp Thr Cys Gln Lys
 20 25 30
 Ser Gly Lys Trp Asn Lys Lys Ser Asn Pro Lys Cys Met Pro Ala Lys
 35 40 45
 Cys Pro Glu Pro Pro Leu Leu Glu Asn Gln Leu Val Leu Lys Glu Leu
 50 55 60
 Thr Thr Glu Val Gly Val Val Thr Phe Ser Cys Lys Glu Gly His Val
 65 70 75 80
 Leu Gln Gly Pro Ser Val Leu Lys Cys Leu Pro Ser Gln Gln Trp Asn
 85 90 95
 Asp Ser Phe Pro Val Cys Lys Ile Val Leu Cys Thr Pro Pro Pro Leu
 100 105 110
 Ile Ser Phe Gly Val Pro Ile Pro Ser Ser Ala Leu His Phe Gly Ser
 115 120 125
 Thr Val Lys Tyr Ser Cys Val Gly Gly Phe Phe Leu Arg Gly Asn Ser
 130 135 140

Thr Thr Leu Cys Gln Pro Asp Gly Thr Trp Ser Ser Pro Leu Pro Glu
 145 150 155 160
 Cys Val Pro Val Glu Cys Pro Gln Pro Glu Glu Ile Pro Asn Gly Ile
 165 170 175
 Ile Asp Val Gln Gly Leu Ala Tyr Leu Ser Thr Ala Leu Tyr Thr Cys
 180 185 190
 Lys Pro Gly Phe Glu Leu Val Gly Asn Thr Thr Thr Leu Cys Gly Glu
 195 200 205
 Asn Gly His Trp Leu Gly Gly Lys Pro Thr Cys Lys Ala Ile Glu Cys
 210 215 220
 Leu Lys Pro Lys Glu Ile Leu Asn Gly Lys Phe Ser Tyr Thr Asp Leu
 225 230 235 240
 His Tyr Gly Gln Thr Val Thr Tyr Ser Cys Asn Arg Gly Phe Arg Leu
 245 250 255
 Glu Gly Pro Ser Ala Leu Thr Cys Leu Glu Thr Gly Asp Trp Asp Val
 260 265 270
 Asp Ala Pro Ser Cys Asn Ala Ile His Cys Asp Ser Pro Gln Pro Ile
 275 280 285
 Glu Asn Gly Phe Val Glu Gly Ala Asp Tyr Ser Tyr Gly Ala Ile Ile
 290 295 300
 Ile Tyr Ser Cys Phe Pro Gly Phe Gln Val Ala Gly His Ala Met Gln
 305 310 315 320
 Thr Cys Glu Glu Ser Gly Trp Ser Ser Ser Ile Pro Thr Cys Met Pro
 325 330 335
 Ile Asp Cys Gly Leu Pro Pro His Ile Asp Phe Gly Ala Cys Thr Lys
 340 345 350
 Leu Lys Asp Ala Arg Asp Ile Leu Ser Lys Lys Arg His Asp Gly Ser
 355 360 365
 Ser Ile Cys Asp Ser Ser Pro Ser Leu Ser Phe Gly Ala Val Ala Lys
 370 375 380
 Thr Trp Glu Asn Thr Lys Glu Ser Pro Ala Thr His Ser Ser Asn Phe
 385 390 395 400
 Leu Tyr Gly Thr Met Val Ser Tyr Thr Cys Asn Pro Gly Tyr Glu Leu
 405 410 415
 Leu Gly Asn Pro Val Leu Ile Cys Gln Glu Asp Gly Thr Trp Asn Gly
 420 425 430
 Ser Ala Pro Ser Cys Ile Ser Ile Glu Cys Asp Leu Pro Thr Ala Pro
 435 440 445
 Glu Asn Gly Phe Leu Arg Phe Thr Glu Thr Ser Met Gly Ser Ala Val
 450 455 460
 Gln Tyr Ser Cys Lys Pro Gly His Ile Leu Ala Gly Ser Asp Leu Arg
 465 470 475 480

Leu

<210> 52
 <211> 200
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Von
 willebrand Factor Type A domain sequence

<400> 52
 Asp Ile Val Phe Leu Leu Asp Gly Ser Gly Ser Ile Gly Ser Gln Asn
 1 5 10 15
 Phe Glu Arg Val Lys Asp Phe Val Glu Arg Val Val Glu Arg Leu Asp
 20 25 30
 Val Gly Pro Arg Asp Lys Lys Glu Glu Asp Ala Val Arg Val Gly Leu
 35 40 45
 Val Gln Tyr Ser Asp Asn Val Arg Thr Glu Ile Lys Phe Lys Leu Asn
 50 55 60
 Asp Tyr Gln Asn Lys Asp Glu Val Leu Gln Ala Leu Gln Lys Ile Arg
 65 70 75 80
 Tyr Glu Asp Tyr Tyr Gly Gly Gly Gly Thr Asn Thr Gly Ala Ala Leu
 85 90 95
 Gln Tyr Val Val Arg Asn Leu Phe Thr Glu Ala Ser Gly Ser Arg Ile
 100 105 110
 Glu Pro Val Ala Glu Glu Gly Ala Pro Lys Val Leu Val Val Leu Thr
 115 120 125
 Asp Gly Arg Ser Gln Asp Asp Pro Ser Pro Thr Ile Asp Ile Arg Asp
 130 135 140
 Val Leu Asn Glu Leu Lys Lys Glu Ala Gly Val Glu Val Phe Ala Ile
 145 150 155 160
 Gly Val Gly Asn Ala Asp Asn Asn Asn Leu Glu Glu Leu Arg Glu Ile
 165 170 175
 Ala Ser Lys Pro Asp Asp His Val Phe Lys Val Ser Asp Phe Glu Ala
 180 185 190
 Leu Asp Thr Leu Gln Glu Leu Leu
 195 200

<210> 53
 <211> 147
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Pentaxin
 domain sequence

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<400> 53

Ser Tyr Ala Thr Lys Lys Pro Leu Lys Asp Asn Glu Leu Leu Ile Phe
 1 5 10 15
 Lys Glu Lys Asp Gly Gln Tyr Ser Leu Tyr Val Gly Gly Ala Pro Gln
 20 25 30
 Leu Glu Val Thr Phe Lys Val Lys Glu Glu Phe Val Ala Pro Val His
 35 40 45
 Ile Cys Thr Ser Trp Glu Ser Ser Ser Gly Ile Ala Glu Phe Trp Val
 50 55 60
 Asp Gly Lys His Cys Pro Trp Val Arg Lys Gly Leu Lys Lys Gly Tyr
 65 70 75 80
 Thr Val Gly Ala Glu Pro Ser Ile Ile Leu Gly Gln Glu Gln Asp Ser
 85 90 95
 Tyr Gly Gly Gly Phe Asp Lys Ser Gln Ser Leu Val Gly Glu Ile Gly
 100 105 110
 Asp Leu Asn Met Trp Asp Tyr Val Leu Thr Pro Glu Glu Ile Lys Thr
 115 120 125
 Val Tyr Lys Gly Ala Gly Pro Leu Glu Arg His Ile Tyr Pro Asn Ile
 130 135 140
 Leu Asp Trp
 145

<210> 54

<211> 62

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Sushi domain sequence

<400> 54

Cys Pro Pro Pro Asp Ile Glu Asn Gly Arg Val Ser Ser Ser Gly Thr
 1 5 10 15
 Tyr Glu Tyr Pro Val Gly Asp Thr Val Thr Tyr Thr Cys Asn Glu Gly
 20 25 30
 Tyr Arg Leu Val Gly Ser Ser Ser Ile Thr Cys Thr Glu Asp Gly Gly
 35 40 45
 Gly Gly Trp Ser Pro Pro Leu Leu Gly Glu Leu Pro Lys Cys
 50 55 60

<210> 55

<211> 207

<212> PRT

<213> Homo sapiens

<400> 55

Met Gly Ser Cys Ser Gly Arg Cys Ala Leu Val Val Leu Cys Ala Phe
 1 5 10 15

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Gln Leu Val Ala Ala Leu Glu Arg Gln Val Phe Asp Phe Leu Gly Tyr
 20 25 30
 Gln Trp Ala Pro Ile Leu Ala Asn Phe Val His Ile Ile Ile Val Ile
 35 40 45
 Leu Gly Leu Phe Gly Thr Ile Gln Tyr Arg Leu Arg Tyr Val Met Tyr
 50 55 60
 Thr Leu Trp Ala Ala Val Trp Val Thr Trp Asn Val Phe Ile Ile Cys
 65 70 75 80
 Phe Tyr Leu Glu Val Gly Gly Leu Leu Lys Asp Ser Glu Leu Leu Thr
 85 90 95
 Phe Ser Leu Ser Arg His Arg Ser Trp Trp Arg Glu Arg Trp Pro Gly
 100 105 110
 Cys Leu His Glu Glu Val Pro Ala Val Gly Leu Gly Ala Pro His Gly
 115 120 125
 Gln Ala Leu Val Ser Gly Ala Gly Cys Ala Leu Glu Pro Ser Tyr Val
 130 135 140
 Glu Ala Leu His Ser Cys Leu Gln Ile Leu Ile Ala Leu Leu Gly Phe
 145 150 155 160
 Val Cys Gly Cys Gln Val Val Ser Val Phe Thr Glu Glu Glu Asp Ser
 165 170 175
 Phe Asp Phe Ile Gly Gly Phe Asp Pro Phe Pro Leu Tyr His Val Asn
 180 185 190
 Glu Lys Pro Ser Ser Leu Leu Ser Lys Gln Val Tyr Leu Pro Ala
 195 200 205

<210> 56
 <211> 208
 <212> PRT
 <213> Mus musculus

<400> 56
 Met Gly Phe Cys Ser Gly Arg Cys Thr Leu Leu Ala Leu Cys Ala Leu
 1 5 10 15
 Gln Leu Val Thr Ala Leu Glu Arg Gln Val Phe Asp Phe Leu Gly Tyr
 20 25 30
 Gln Trp Ala Pro Ile Leu Ala Asn Phe Thr His Ile Ile Val Val Ile
 35 40 45
 Leu Gly Leu Phe Gly Thr Ile Gln Tyr Arg Pro Arg Tyr Ile Val Val
 50 55 60
 Tyr Val Val Trp Ala Ala Val Trp Val Thr Trp Asn Val Phe Ile Ile
 65 70 75 80
 Cys Phe Tyr Leu Glu Val Gly Gly Leu Ser Lys Asp Ser Glu Leu Leu
 85 90 95
 Thr Phe Asn Leu Ser Gly His Arg Ser Trp Trp Glu Glu His Gly Pro
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100 105 110
 Gly Cys Leu His Glu Glu Glu Ala Thr Ala Gly Leu Gly Ala Leu His
 115 120 125
 Gly Gln Ser Leu Val Val Gly Ala Gly Cys Ala Met Val His Ser Tyr
 130 135 140
 Val Glu Ala Leu His Ser Gly Leu Gln Ile Leu Leu Ala Leu Leu Gly
 145 150 155 160
 Phe Val Tyr Gly Cys Tyr Val Val Ser Val Leu Thr Glu Glu Glu Asp
 165 170 175
 Ser Phe Asp Phe Ile Gly Gly Phe Asp Pro Phe Pro Leu Tyr His Val
 180 185 190
 Asn Glu Lys Pro Ser Ser Leu Leu Ser Lys Gln Ala Tyr Leu Pro Ala
 195 200 205

<210> 57
 <211> 208
 <212> PRT
 <213> Mus musculus

<400> 57
 Met Gly Phe Cys Ser Gly Arg Cys Thr Leu Leu Ala Leu Cys Arg Leu
 1 5 10 15
 Gln Leu Val Thr Ala Leu Glu Arg Gln Val Phe Asp Phe Leu Gly Tyr
 20 25 30
 Gln Trp Ala Pro Ile Leu Ala Asn Phe Thr His Ile Ile Val Val Ile
 35 40 45
 Leu Gly Leu Phe Gly Thr Ile Gln Tyr Arg Pro Arg Tyr Ile Val Val
 50 55 60
 Tyr Val Val Trp Ala Ala Val Trp Val Thr Trp Asn Val Phe Ile Ile
 65 70 75 80
 Cys Phe Tyr Leu Glu Val Gly Gly Leu Ser Lys Asp Ser Glu Leu Leu
 85 90 95
 Thr Phe Asn Leu Ser Gly His Arg Ser Trp Trp Glu Glu His Gly Pro
 100 105 110
 Gly Cys Leu His Glu Glu Glu Ala Thr Ala Gly Leu Gly Ala Leu His
 115 120 125
 Gly Gln Ser Leu Val Val Gly Ala Gly Cys Ala Met Val His Ser Tyr
 130 135 140
 Val Glu Ala Leu His Ser Gly Leu Gln Ile Leu Leu Ala Leu Leu Gly
 145 150 155 160
 Phe Val Tyr Gly Cys Tyr Val Val Ser Val Leu Thr Glu Glu Glu Asp
 165 170 175

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Ser Phe Asp Phe Ile Gly Gly Phe Asp Pro Phe Pro Leu Tyr His Val
 180 185 190

Asn Glu Lys Pro Ser Ser Leu Leu Ser Lys Gln Ala Tyr Leu Pro Ala
 195 200 205

<210> 58
 <211> 208
 <212> PRT
 <213> Mus musculus

<400> 58
 Met Gly Phe Cys Ser Gly Arg Cys Thr Leu Leu Ala Leu Cys Ala Leu
 1 5 10 15

Gln Leu Val Thr Ala Leu Glu Arg Gln Val Phe Asp Phe Leu Gly Tyr
 20 25 30

Gln Trp Ala Pro Ile Leu Ala Asn Phe Thr His Ile Ile Val Val Ile
 35 40 45

Leu Gly Leu Phe Gly Thr Ile Gln Tyr Arg Pro Arg Tyr Ile Val Val
 50 55 60

Tyr Val Val Trp Ala Ala Val Trp Val Thr Trp Asn Val Phe Ile Ile
 65 70 75 80

Cys Phe Tyr Leu Glu Val Gly Gly Leu Ser Lys Asp Ser Glu Leu Leu
 85 90 95

Thr Phe Asn Leu Ser Gly His Arg Ser Trp Trp Glu Glu His Gly Pro
 100 105 110

Gly Cys Leu His Glu Glu Glu Ala Thr Ala Gly Leu Gly Ala Leu His
 115 120 125

Gly Gln Ser Leu Val Val Gly Ala Gly Cys Ala Met Val His Ser Tyr
 130 135 140

Val Glu Ala Leu His Ser Gly Leu Gln Ile Leu Leu Ala Leu Leu Gly
 145 150 155 160

Phe Val Tyr Gly Cys Tyr Val Val Arg Val Leu Thr Glu Glu Glu Asp
 165 170 175

Ser Phe Asp Phe Ile Gly Gly Phe Asp Pro Phe Pro Leu Tyr His Val
 180 185 190

Asn Glu Lys Pro Ser Ser Leu Leu Ser Lys Gln Ala Tyr Leu Pro Ala
 195 200 205

<210> 59
 <211> 207
 <212> PRT
 <213> Mus musculus

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<400> 59

Met Gly Lys Cys Ser₅ Gly Arg Cys Thr Leu₁₀ Val Ala Phe Cys Cys₁₅ Leu
 Gln Leu Val Ala₂₀ Ala Leu Gln Arg Gln₂₅ Ile Phe Asp Phe Leu₃₀ Gly Tyr
 Gln Trp Ala₃₅ Pro Ile Leu Ala Asn₄₀ Phe Leu His Ile Met₄₅ Ala Val Ile
 Leu Gly Ile Phe Gly Thr Val₅₅ Gln Tyr Arg Ser Arg₆₀ Tyr Leu Ile Leu
 Tyr Ala Ala Trp Leu Val₇₀ Leu Trp Val Gly Trp₇₅ Asn Ala Phe Ile Ile₈₀
 Cys Phe Tyr Leu Glu₈₅ Val Gly Gln Leu Ser₉₀ Gln Asp Arg Asp Phe₉₅ Ile
 Met Thr Phe Asn₁₀₀ Thr Ser Leu His Arg₁₀₅ Ser Trp Trp Met Glu₁₁₀ Asn Gly
 Pro Gly Cys₁₁₅ Leu Val Thr Pro Val₁₂₀ Leu Asn Ser Arg Leu₁₂₅ Ala Leu Glu
 Asp His His Val Ile Ser Val₁₃₅ Thr Gly Cys Leu Leu₁₄₀ Asp Tyr Pro Tyr
 Ile Glu Ala Leu Ser₁₅₀ Ser Ala Leu Gln Ile Phe₁₅₅ Leu Ala Leu Phe Gly₁₆₀
 Phe Val Phe Ala Cys₁₆₅ Tyr Val Ser Lys Val₁₇₀ Phe Leu Glu Glu Glu₁₇₅ Asp
 Ser Phe Asp Phe₁₈₀ Ile Gly Gly Phe Asp₁₈₅ Ser Tyr Gly Tyr Gln₁₉₀ Ala Pro
 Gln Lys Thr₁₉₅ Ser His Leu Gln Leu₂₀₀ Gln Pro Leu Tyr Thr₂₀₅ Ser Gly

<210> 60

<211> 367

<212> PRT

<213> Mus musculus

<400> 60

Met Trp Gly Ser Arg₅ Ala Gln Gln Ser Gly₁₀ Pro Asp Arg Gly Gly₁₅ Ala
 Cys Leu Leu Ala₂₀ Ala Phe Leu Leu Cys₂₅ Phe Ser Leu Leu His₃₀ Ala Gln
 Asp Tyr Thr₃₅ Pro Ser Gln Thr Pro₄₀ Pro Pro Thr Ser Asn₄₅ Thr Ser Leu
 Lys Pro Arg Gly Arg Val Gln₅₅ Lys Glu Leu Cys Gly₆₀ Lys Thr Lys Phe
 Gln Gly Lys Ile Tyr Gly₇₀ Gly Gln Ile Ala Lys₇₅ Ala Glu Arg Trp Pro₈₀

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Trp Gln Ala Ser Leu Ile Phe Arg Gly Arg His Ile Cys Gly Ala Val
85 90 95

Leu Ile Asp Lys Thr Trp Leu Leu Ser Ala Ala His Cys Phe Gln Arg
100 105 110

Ser Leu Thr Pro Ser Asp Tyr Arg Ile Leu Leu Gly Tyr Asn Gln Leu
115 120 125

Ser Asn Pro Ser Asn Tyr Ser Arg Gln Met Thr Val Asn Lys Val Ile
130 135 140

Leu His Glu Asp Tyr Ser Lys Leu Ser Arg Leu Glu Lys Asn Ile Val
145 150 155 160

Leu Ile Gln Leu His His Pro Val Ile Tyr Ser Thr His Ile Phe Pro
165 170 175

Ala Cys Val Pro Asp Gly Thr Thr Lys Val Ser Pro Asn Asn Leu Cys
180 185 190

Trp Ile Ser Gly Trp Gly Met Leu Ser Ala Asp Lys Phe Leu Gln Ala
195 200 205

Pro Phe Pro Leu Leu Asp Ala Glu Val Ser Leu Ile Asp Glu Glu Glu
210 215 220

Cys Thr Thr Phe Phe Gln Thr Pro Glu Val Ser Ile Thr Glu Tyr Asp
225 230 235 240

Val Ile Lys Asp Asp Val Leu Cys Ala Gly Asp Leu Thr Asn Gln Lys
245 250 255

Ser Ser Cys Arg Gly Asp Ser Gly Gly Pro Leu Val Cys Phe Leu Asn
260 265 270

Ser Phe Trp Tyr Val Val Gly Leu Ala Asn Trp Asn Gly Ala Cys Leu
275 280 285

Glu Pro Ile His Ser Pro Asn Ile Phe Thr Lys Val Ser Tyr Phe Ser
290 295 300

Asp Trp Ile Lys Gln Lys Lys Ala Asn Thr Pro Ala Ala Asp Val Ser
305 310 315 320

Ser Ala Pro Leu Glu Glu Met Ala Ser Ser Leu Arg Gly Trp Gly Asn
325 330 335

Tyr Ser Ala Gly Ile Thr Leu Lys Pro Arg Ile Ser Thr Thr Leu Leu
340 345 350

Ser Ser Gln Ala Leu Leu Leu Gln Ser Ile Trp Leu Arg Ile Leu
355 360 365

<210> 61
<211> 366
<212> PRT
<213> Mus musculus

<400> 61
Met Cys Gly Val Arg Ala Lys Lys Ser Gly Leu Ser Gly Tyr Gly Ala
1 5 10 15

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Gly Leu Leu Ala Ala Leu Leu Gly Val Ser Phe Leu Ser Gln His Ala
 20 25 30
 Gln Thr Ala Glu Pro Thr Asn Val Thr Asn Ala Ala Asn Asn Thr Thr
 35 40 45
 Ile Gln Ile Met Lys Ser Thr Leu Ser Leu Ser Glu Val Cys Gly Lys
 50 55 60
 Thr Lys Phe Gln Gly Lys Ile Tyr Gly Gly Gln Ile Ala Gly Ala Glu
 65 70 75 80
 Arg Trp Pro Trp Gln Ala Ser Leu Arg Leu Tyr Gly Arg His Ile Cys
 85 90 95
 Gly Ala Val Leu Ile Asp Lys Asn Trp Val Leu Gly Ala Ala His Cys
 100 105 110
 Phe Gln Arg Ser Gln Glu Pro Ser Asp Tyr His Val Met Leu Gly Tyr
 115 120 125
 Thr Asp Leu Asn Ser Pro Thr Arg Tyr Ser Arg Thr Met Ser Val Gln
 130 135 140
 Lys Val Ile Val His Lys Asp Tyr Asn Arg Phe His Thr Gln Gly Ser
 145 150 155 160
 Asp Ile Val Leu Leu Gln Leu Arg Ser Ser Val Glu Tyr Ser Ser His
 165 170 175
 Ile Leu Pro Ala Cys Val Pro Glu Glu Asn Ile Lys Ile Pro Lys Glu
 180 185 190
 Lys Ala Cys Trp Ala Ser Gly Trp Gly Tyr Leu Arg Glu Asp Val Arg
 195 200 205
 Ile Pro Leu Pro Asn Glu Leu Tyr Glu Ala Glu Leu Ile Ile Met Ser
 210 215 220
 Asn Asp Gln Cys Lys Gly Phe Phe Pro Pro Pro Val Pro Gly Ser Ser
 225 230 235 240
 Arg Ser Tyr Tyr Ile Tyr Asp Asp Met Val Cys Ala Ala Asp Tyr Asp
 245 250 255
 Met Ser Lys Ser Ile Cys Ala Gly Asp Ser Gly Gly Pro Leu Val Cys
 260 265 270
 Leu Leu Glu Gly Ser Trp Tyr Val Val Gly Leu Thr Ser Trp Ser Ser
 275 280 285
 Thr Cys Glu Glu Pro Ile Val Ser Pro Ser Val Phe Ala Arg Val Ser
 290 295 300
 Tyr Phe Asp Lys Trp Ile Lys Asp Asn Lys Lys Ser Ser Ser Asn Ser
 305 310 315 320
 Lys Pro Gly Glu Ser Pro His His Pro Gly Ser Pro Glu Asn Glu Asn
 325 330 335
 Pro Glu Gly Asn Asn Lys Asn Gln Gly Thr Val Ile Lys Pro Val Cys
 340 345 350

Thr Ala Leu Leu Leu Ser Gln Thr Leu Leu Gln Gln Leu Ile
 355 360 365

<210> 62
 <211> 143
 <212> PRT
 <213> Mus musculus

<400> 62
 Met Leu Gly Tyr Thr 5 Asp Leu Asn Ser Pro Thr Arg Tyr Ser Arg Thr
 1 10 15
 Met Ser Val Gln Lys Val Ile Val His Lys Asp Tyr Asn Arg Phe His
 20 25 30
 Thr Gln Gly Ser Asp Ile Val Leu Leu Gln Leu Arg Ser Ser Val Glu
 35 40 45
 Tyr Ser Ser His Ile Leu Pro Ala Cys Val Pro Glu Glu Asn Ile Lys
 50 55 60
 Ile Pro Lys Glu Lys Ala Cys Trp Ala Ser Gly Trp Gly Tyr Leu Arg
 65 70 75 80
 Glu Asp Val Arg Ile Pro Leu Pro Asn Glu Leu Tyr Glu Ala Glu Leu
 85 90 95
 Ile Ile Met Ser Asn Asp Gln Cys Lys Gly Phe Phe Pro Pro Pro Val
 100 105 110
 Pro Gly Ser Gly Arg Ser Tyr Tyr Ile Tyr Asp Asp Met Val Cys Ala
 115 120 125
 Ala Asp Tyr Asp Met Ser Lys Ser Ile Cys Ala Gly Leu Leu Leu
 130 135 140

<210> 63
 <211> 273
 <212> PRT
 <213> Ovis aries

<400> 63
 Met Leu His Leu Leu Ala Leu Ala Leu Leu Leu Ser Leu Val Ser Ala
 1 5 10 15
 Ala Pro Ala Pro Gly Gln Ala Leu Gln Arg Ser Gly Ile Ile Gly Gly
 20 25 30
 Lys Glu Ala Pro Gly Ser Arg Trp Pro Trp Gln Val Ser Leu Arg Val
 35 40 45
 Arg Asp Gln Tyr Trp Arg His Gln Cys Gly Gly Ser Leu Ile His Pro
 50 55 60
 Gln Trp Val Leu Thr Ala Ala His Cys Ile Gly Pro Glu Leu Gln Glu
 65 70 75 80
 Pro Ser Asp Phe Arg Val Gln Leu Arg Glu Gln His Leu Tyr Tyr Gln
 85 90 95

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Asp Arg Leu Leu Pro Ile Ser Arg Val Ile Pro His Pro His Tyr Tyr
100 105 110

Met Val Glu Asn Gly Ala Asp Ile Ala Leu Leu Gln Leu Glu Glu Pro
115 120 125

Val Ser Ile Ser Arg His Val Gln Pro Val Thr Leu Pro Pro Ala Ser
130 135 140

Glu Thr Phe Pro Pro Glu Ser Gln Cys Trp Val Thr Gly Trp Gly Asp
145 150 155 160

Val Asp Asn Gly Arg Pro Leu Pro Pro Pro Tyr Pro Leu Lys Gln Val
165 170 175

Lys Val Pro Ile Val Glu Asn Ser Val Cys Asp Trp Lys Tyr His Ser
180 185 190

Gly Leu Ser Thr Asp Tyr Ser Val Pro Ile Val Gln Glu Asp Asn Leu
195 200 205

Cys Ala Gly Asp Gly Gly Arg Asp Ser Cys Gln Gly Asp Ser Gly Gly
210 215 220

Pro Leu Val Cys Lys Val Asn Gly Thr Trp Leu Gln Ala Gly Val Val
225 230 235 240

Ser Trp Gly Asp Gly Cys Ala Lys Pro Asn Arg Pro Gly Ile Tyr Thr
245 250 255

Arg Ile Thr Ser Tyr Leu Asp Trp Ile His Gln Tyr Val Pro Gln Glu
260 265 270

Pro

<210> 64
<211> 273
<212> PRT
<213> Ovis aries

<400> 64
Met Leu His Leu Leu Ala Leu Ala Leu Leu Leu Ser Leu Val Ser Ala
1 5 10 15

Ala Pro Gly Pro Gly Gln Ala Leu Gln Arg Ser Gly Ile Ile Gly Gly
20 25 30

Lys Glu Ala Pro Gly Ser Arg Trp Pro Trp Gln Val Ser Leu Arg Val
35 40 45

Arg Asp Gln Tyr Trp Arg His Gln Cys Gly Gly Ser Leu Ile His Pro
50 55 60

Gln Trp Val Leu Thr Ala Ala His Cys Ile Gly Pro Glu Leu Gln Glu
65 70 75 80

Pro Ser Asp Phe Arg Val Gln Leu Arg Glu Gln His Leu Tyr Tyr Gln
85 90 95

Asp Arg Leu Leu Pro Ile Ser Arg Val Ile Pro His Pro His Tyr Tyr
100 105 110

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Met Val Glu Asn Gly Ala Asp Ile Ala Leu Leu Gln Leu Glu Glu Pro
 115 120 125
 Val Ser Ile Ser Cys His Val Arg Pro Val Thr Leu Pro Pro Ala Ser
 130 135 140
 Glu Thr Phe Pro Pro Gly Ser Gln Cys Trp Val Thr Gly Trp Gly Asn
 145 150 155 160
 Val Asp Asn Gly Arg Pro Leu Pro Pro Pro Tyr Pro Leu Lys Gln Val
 165 170 175
 Lys Val Pro Ile Val Glu Asn Ser Val Cys Asp Trp Lys Tyr His Ser
 180 185 190
 Gly Leu Ser Thr Asp Tyr Ser Val Pro Ile Val Gln Glu Asp Asn Leu
 195 200 205
 Cys Ala Gly Asp Gly Gly Arg Asp Ser Cys Gln Gly Asp Ser Gly Gly
 210 215 220
 Pro Leu Val Cys Lys Val Asn Gly Thr Trp Leu Gln Ala Gly Val Val
 225 230 235 240
 Ser Trp Gly Asp Gly Cys Ala Asn Pro Asp Tyr Pro Gly Val Tyr Thr
 245 250 255
 Arg Ile Thr Ser Tyr Leu Asp Trp Ile His Gln Tyr Val Pro Gln Glu
 260 265 270
 Pro

<210> 65
 <211> 205
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Trypsin
 Domian sequence

<400> 65
 Ser Ala Pro Ala Ser Ser Val Arg Val Ser Leu Ser Val Arg Leu Gly
 1 5 10 15
 Glu His Asn Leu Ser Leu Thr Glu Gly Thr Glu Gln Lys Phe Asp Val
 20 25 30
 Lys Lys Thr Ile Ile Val His Pro Asn Tyr Asn Pro Asp Thr Leu Asp
 35 40 45
 Asn Gly Ala Tyr Asp Asn Asp Ile Ala Leu Leu Lys Leu Lys Ser Pro
 50 55 60
 Gly Val Thr Leu Gly Asp Thr Val Arg Pro Ile Cys Leu Pro Ser Ala
 65 70 75 80
 Ser Ser Asp Leu Pro Val Gly Thr Thr Cys Thr Val Ser Gly Trp Gly
 85 90 95

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Arg	Arg	Pro	Thr	Lys	Asn	Leu	Gly	Leu	Ser	Asp	Thr	Leu	Gln	Glu	Val
			100					105					110		
Val	Val	Pro	Val	Val	Ser	Arg	Glu	Thr	Cys	Arg	Ser	Ala	Tyr	Glu	Tyr
		115					120					125			
Gly	Gly	Thr	Asp	Asp	Lys	Val	Glu	Phe	Val	Thr	Asp	Asn	Met	Ile	Cys
	130					135					140				
Ala	Gly	Ala	Leu	Gly	Gly	Lys	Asp	Ala	Cys	Gln	Gly	Asp	Ser	Gly	Gly
145					150					155					160
Pro	Leu	Val	Cys	Ser	Asp	Gly	Asn	Arg	Asp	Gly	Arg	Trp	Glu	Leu	Val
			165						170					175	
Gly	Ile	Val	Ser	Trp	Gly	Ser	Tyr	Gly	Cys	Ala	Arg	Gly	Asn	Lys	Pro
			180					185					190		
Gly	Val	Tyr	Thr	Arg	Val	Ser	Ser	Tyr	Leu	Asp	Trp	Ile			
		195					200					205			

<210> 66
 <211> 349
 <212> PRT
 <213> Homo sapiens

<400> 66

Met	Asn	Arg	Lys	Ala	Leu	Arg	Cys	Leu	Gly	His	Leu	Phe	Leu	Ser	Leu
1				5					10					15	
Gly	Met	Val	Cys	Leu	Arg	Ile	Gly	Gly	Phe	Ser	Ser	Val	Val	Ala	Leu
			20					25					30		
Gly	Ala	Thr	Ile	Ile	Cys	Asn	Lys	Ile	Pro	Gly	Leu	Ala	Pro	Arg	Gln
		35					40					45			
Arg	Ala	Ile	Cys	Gln	Ser	Arg	Pro	Asp	Ala	Ile	Ile	Val	Ile	Gly	Glu
	50					55					60				
Gly	Ser	Gln	Met	Gly	Leu	Asp	Glu	Cys	Gln	Phe	Gln	Phe	Arg	Asn	Gly
65					70				75					80	
Arg	Trp	Asn	Cys	Ser	Ala	Leu	Gly	Glu	Arg	Thr	Val	Phe	Gly	Lys	Glu
			85						90					95	
Leu	Lys	Val	Gly	Ser	Arg	Asp	Gly	Ala	Phe	Thr	Tyr	Ala	Ile	Ile	Ala
			100					105					110		
Ala	Gly	Val	Ala	His	Ala	Ile	Thr	Ala	Ala	Cys	Thr	His	Gly	Asn	Leu
		115					120					125			
Ser	Asp	Cys	Gly	Cys	Asp	Lys	Glu	Lys	Gln	Gly	Gln	Tyr	His	Arg	Asp
	130					135					140				
Glu	Gly	Trp	Lys	Trp	Gly	Gly	Cys	Ser	Ala	Asp	Ile	Arg	Tyr	Gly	Ile
145					150					155					160
Gly	Phe	Ala	Lys	Val	Phe	Val	Asp	Ala	Arg	Glu	Ile	Lys	Gln	Asn	Ala
			165						170					175	
Arg	Thr	Leu	Met	Asn	Leu	His	Asn	Asn	Glu	Ala	Gly	Arg	Lys	Ile	Leu
			180					185					190		

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Glu Glu Asn Met Lys Leu Glu Cys Lys Cys His Gly Val Ser Gly Ser
 195 200 205
 Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Gln Phe Arg Glu Leu
 210 215 220
 Gly Tyr Val Leu Lys Asp Lys Tyr Asn Glu Ala Val His Val Glu Pro
 225 230 235 240
 Val Arg Ala Ser Arg Asn Lys Arg Pro Thr Phe Leu Lys Ile Lys Lys
 245 250 255
 Pro Leu Ser Tyr Arg Lys Pro Met Asp Thr Asp Leu Val Tyr Ile Glu
 260 265 270
 Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly Ser Val Gly
 275 280 285
 Thr Gln Gly Arg Ala Cys Asn Lys Thr Ala Pro Gln Ala Ser Gly Cys
 290 295 300
 Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln Tyr Ala Arg
 305 310 315 320
 Val Trp Gln Cys Asn Cys Lys Phe His Trp Cys Cys Tyr Val Lys Cys
 325 330 335
 Asn Thr Cys Ser Glu Arg Thr Glu Met Tyr Thr Cys Lys
 340 345

<210> 67
 <211> 349
 <212> PRT
 <213> Homo sapiens

<400> 67
 Met Asn Arg Lys Ala Arg Arg Cys Leu Gly His Leu Phe Leu Ser Leu
 1 5 10 15
 Gly Met Val Tyr Leu Arg Ile Gly Gly Phe Ser Ser Val Val Ala Leu
 20 25 30
 Gly Ala Ser Ile Ile Cys Asn Lys Ile Pro Gly Leu Ala Pro Arg Gln
 35 40 45
 Arg Ala Ile Cys Gln Ser Arg Pro Asp Ala Ile Ile Val Ile Gly Glu
 50 55 60
 Gly Ser Gln Met Gly Leu Asp Glu Cys Gln Phe Gln Phe Arg Asn Gly
 65 70 75 80
 Arg Trp Asn Cys Ser Ala Leu Gly Glu Arg Thr Val Phe Gly Lys Glu
 85 90 95
 Leu Lys Val Gly Ser Arg Glu Ala Ala Phe Thr Tyr Ala Ile Ile Ala
 100 105 110
 Ala Gly Val Ala His Ala Ile Thr Ala Ala Cys Thr Gln Gly Asn Leu
 115 120 125
 Ser Asp Cys Gly Cys Asp Lys Glu Lys Gln Gly Gln Tyr His Arg Asp
 Page 146

130 135 140
 Glu Gly Trp Lys Trp Gly Gly Cys Ser Ala Asp Ile Arg Tyr Gly Ile
 145 150 155 160
 Gly Phe Ala Lys Val Phe Val Asp Ala Arg Glu Ile Lys Gln Asn Ala
 165 170 175
 Arg Thr Leu Met Asn Leu His Asn Asn Glu Ala Gly Arg Lys Ile Leu
 180 185 190
 Glu Glu Asn Met Lys Leu Glu Cys Lys Cys His Gly Val Ser Gly Ser
 195 200 205
 Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Gln Phe Arg Glu Leu
 210 215 220
 Gly Tyr Val Leu Lys Asp Lys Tyr Asn Glu Ala Val His Val Glu Pro
 225 230 235 240
 Val Arg Ala Ser Arg Asn Lys Arg Pro Thr Phe Leu Lys Ile Lys Lys
 245 250 255
 Pro Leu Ser Tyr Arg Lys Pro Met Asp Thr Asp Leu Val Tyr Ile Glu
 260 265 270
 Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly Ser Val Gly
 275 280 285
 Thr Gln Gly Arg Ala Cys Asn Lys Thr Ala Pro Gln Ala Ser Gly Cys
 290 295 300
 Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln Tyr Ala Arg
 305 310 315 320
 Val Trp Gln Cys Asn Cys Lys Phe His Trp Cys Cys Tyr Val Lys Cys
 325 330 335
 Asn Thr Cys Ser Glu Arg Thr Glu Met Tyr Thr Cys Lys
 340 345

<210> 68
 <211> 349
 <212> PRT
 <213> Mus musculus

<400> 68
 Met Thr Arg Lys Ala Arg Arg Cys Leu Gly His Leu Phe Leu Ser Leu
 1 5 10 15
 Gly Ile Val Tyr Leu Arg Ile Gly Gly Phe Ser Ser Val Val Ala Leu
 20 25 30
 Gly Ala Ser Ile Ile Cys Asn Lys Ile Pro Gly Leu Ala Pro Arg Gln
 35 40 45
 Arg Ala Ile Cys Gln Ser Arg Pro Asp Ala Ile Ile Val Ile Gly Glu
 50 55 60
 Gly Ser Gln Met Gly Leu Asp Glu Cys Gln Phe Gln Phe Arg Asn Gly
 65 70 75 80

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Arg Trp Asn Cys Ser Ala Leu Gly Glu Arg Thr Val Phe Gly Lys Glu
 85 90 95
 Leu Lys Val Gly Ser Arg Glu Ala Ala Phe Thr Tyr Ala Ile Ile Ala
 100 105 110
 Ala Gly Val Ala His Ala Ile Thr Ala Ala Cys Thr Gln Gly Asn Leu
 115 120 125
 Ser Asp Cys Gly Cys Asp Lys Glu Lys Gln Gly Gln Tyr His Arg Asp
 130 135 140
 Glu Gly Trp Lys Trp Gly Gly Cys Ser Ala Asp Ile Arg Tyr Gly Ile
 145 150 155 160
 Gly Phe Ala Lys Val Phe Val Asp Ala Arg Glu Ile Lys Gln Asn Ala
 165 170 175
 Arg Thr Leu Met Asn Leu His Asn Asn Glu Ala Gly Arg Lys Ile Leu
 180 185 190
 Glu Glu Asn Met Lys Leu Glu Cys Lys Cys His Gly Val Ser Gly Ser
 195 200 205
 Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Gln Phe Arg Glu Leu
 210 215 220
 Gly Tyr Val Leu Lys Asp Lys Tyr Asn Glu Ala Val His Val Glu Pro
 225 230 235 240
 Val Arg Ala Ser Arg Asn Lys Arg Pro Thr Phe Leu Lys Ile Lys Lys
 245 250 255
 Pro Leu Ser Tyr Arg Lys Pro Met Asp Thr Asp Leu Val Tyr Ile Glu
 260 265 270
 Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly Ser Val Gly
 275 280 285
 Thr Gln Gly Arg Ala Cys Asn Lys Thr Ala Pro Gln Ala Ser Gly Cys
 290 295 300
 Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln Tyr Ala Arg
 305 310 315 320
 Val Trp Gln Cys Asn Cys Lys Phe His Trp Cys Cys Tyr Val Lys Cys
 325 330 335
 Asn Thr Cys Ser Glu Arg Thr Glu Met Tyr Thr Cys Lys
 340 345

<210> 69
 <211> 349
 <212> PRT
 <213> Mus musculus

<400> 69
 Met Thr Arg Lys Ala Arg Arg Cys Leu Gly His Leu Phe Leu Ser Leu
 1 5 10 15
 Gly Ile Val Tyr Leu Arg Ile Gly Gly Phe Ser Ser Val Val Ala Leu
 20 25 30

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Gly Ala Ser₃₅ Ile Ile Cys Asn Lys₄₀ Ile Pro Gly Leu Ala₄₅ Pro Arg Gln
 Arg Ala₅₀ Ile Cys Gln Ser Arg₅₅ Pro Asp Ala Ile Ile₆₀ Val Ile Gly Glu
 Gly₆₅ Ser Gln Met Gly Leu₇₀ Asp Glu Cys Gln Phe₇₅ Gln Phe Arg Asn Gly₈₀
 Arg Trp Asn Cys Ser₈₅ Ala Leu Gly Glu Arg₉₀ Thr Val Phe Gly Lys₉₅ Glu
 Leu Lys Val Gly₁₀₀ Ser Arg Glu Ala₁₀₅ Phe Thr Tyr Ala₁₁₀ Ile Ile Ala
 Ala Gly Val₁₁₅ Ala His Ala Ile Thr₁₂₀ Ala Ala Cys Thr Gln₁₂₅ Gly Asn Leu
 Ser Asp₁₃₀ Cys Gly Cys Asp Lys₁₃₅ Glu Lys Gln Gly Gln₁₄₀ Tyr His Trp Asp
 Glu₁₄₅ Gly Trp Lys Trp Gly₁₅₀ Gly Cys Ser Ala Asp₁₅₅ Ile Arg Tyr Gly Ile₁₆₀
 Gly Phe Ala Lys Val₁₆₅ Phe Val Asp Ala Arg₁₇₀ Glu Ile Lys Gln Asn₁₇₅ Ala
 Arg Thr Leu Met₁₈₀ Asn Leu His Asn₁₈₅ Glu Ala Gly Arg Lys₁₉₀ Ile Leu
 Glu Glu Asn₁₉₅ Met Lys Leu Glu Cys₂₀₀ Lys Cys His Gly Val₂₀₅ Ser Gly Ser
 Cys Thr₂₁₀ Thr Lys Thr Cys Trp₂₁₅ Thr Thr Leu Pro Gln₂₂₀ Phe Arg Glu Leu
 Gly₂₂₅ Tyr Val Leu Lys Asp₂₃₀ Lys Tyr Asn Glu Ala₂₃₅ Val His Val Glu Pro₂₄₀
 Val Arg Ala Ser Arg₂₄₅ Asn Lys Arg Pro Thr₂₅₀ Phe Leu Lys Ile Lys₂₅₅ Lys
 Pro Leu Ser Tyr₂₆₀ Arg Lys Pro Met Asp₂₆₅ Thr Asp Leu Val Tyr₂₇₀ Ile Glu
 Leu Ser Pro₂₇₅ Asn Tyr Cys Glu Glu₂₈₀ Asp Pro Val Thr Gly₂₈₅ Ser Val Gly
 Thr Gln₂₉₀ Gly Arg Ala Cys Asn₂₉₅ Lys Thr Ala Pro Gln₃₀₀ Ala Ser Gly Cys
 Asp₃₀₅ Leu Met Cys Cys Gly₃₁₀ Arg Gly Tyr Asn Thr₃₁₅ His Gln Tyr Ala Arg₃₂₀
 Val Trp Gln Cys Asn₃₂₅ Cys Lys Phe His Trp₃₃₀ Cys Cys Tyr Val Lys₃₃₅ Cys
 Asn Thr Cys Ser₃₄₀ Glu Arg Thr Glu Met₃₄₅ Tyr Thr Cys Lys

<210> 70

<211> 349
 <212> PRT
 <213> Gallus gallus

<400> 70

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Met Asn Arg Lys Thr Arg Arg Trp Ile Phe His Ile Phe Leu Ser Leu
 1          5          10          15
Gly Ile Val Tyr Ile Lys Ile Gly Gly Phe Ser Ser Val Val Ala Leu
          20          25          30
Gly Ala Ser Ile Ile Cys Asn Lys Ile Pro Gly Leu Ala Pro Arg Gln
          35          40          45
Arg Ala Ile Cys Gln Ser Arg Pro Asp Ala Ile Ile Val Ile Gly Glu
          50          55          60
Gly Ser Gln Met Gly Ile Asn Glu Cys Gln Phe Gln Phe Arg Asn Gly
          65          70          75          80
Arg Trp Asn Cys Ser Ala Leu Gly Glu Arg Thr Val Phe Gly Lys Glu
          85          90          95
Leu Lys Val Gly Ser Arg Glu Ala Ala Phe Thr Tyr Ala Ile Ile Ala
          100          105          110
Ala Gly Val Ala His Ala Ile Thr Ala Ala Cys Thr Gln Gly Asn Leu
          115          120          125
Ser Asp Cys Gly Cys Asp Lys Glu Lys Gln Gly Gln Tyr His Lys Glu
          130          135          140
Glu Gly Trp Lys Trp Gly Gly Cys Ser Ala Asp Ile Arg Tyr Gly Ile
          145          150          155          160
Gly Phe Ala Lys Val Phe Val Asp Ala Arg Glu Ile Lys Gln Asn Ala
          165          170          175
Arg Thr Leu Met Asn Leu His Asn Asn Glu Ala Gly Arg Lys Ile Leu
          180          185          190
Glu Glu Asn Met Lys Leu Glu Cys Lys Cys His Gly Val Ser Gly Ser
          195          200          205
Cys Thr Thr Lys Thr Cys Trp Thr Thr Leu Pro Lys Phe Arg Glu Leu
          210          215          220
Gly Tyr Ile Leu Lys Asp Lys Tyr Asn Glu Ala Val Gln Val Glu Pro
          225          230          235          240
Val Arg Ala Ser Arg Asn Lys Arg Pro Thr Phe Leu Lys Ile Lys Lys
          245          250          255
Pro Leu Ser Tyr Arg Lys Pro Met Asp Thr Asp Leu Val Tyr Ile Glu
          260          265          270
Lys Ser Pro Asn Tyr Cys Glu Glu Asp Pro Val Thr Gly Ser Val Gly
          275          280          285
Thr Gln Gly Arg Met Cys Asn Lys Thr Ala Gln Gln Ser Asn Gly Cys
          290          295          300
Asp Leu Met Cys Cys Gly Arg Gly Tyr Asn Thr His Gln Tyr Ser Arg

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305 310 320
Val Trp Gln Cys Asn Cys Lys Phe His Trp Cys Cys Tyr Val Lys Cys
 325 330 335
Asn Thr Cys Ser Glu Arg Thr Glu Val Tyr Thr Cys Lys
 340 345

<210> 71
<211> 352
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: WNT domain
sequence

<400> 71
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Arg Arg Asn Pro Asp Val Met Ala Ser Val Ser Glu Gly Ala Gln Leu
 20 25 30
Ala Ile Gln Glu Cys Gln His Gln Phe Arg Gly Arg Arg Trp Asn Cys
 35 40 45
Ser Thr Leu Asp Ser Leu Asn Glu Arg Ser Val Phe Gly Lys Val Leu
 50 55 60
Lys Lys Gly Thr Arg Glu Thr Ala Phe Val Tyr Ala Ile Ser Ser Ala
65 70 75 80
Gly Val Ala His Ala Val Thr Arg Ala Cys Ser Glu Gly Glu Leu Glu
 85 90 95
Ser Cys Gly Cys Asp Asp Lys Arg Lys Ala Asp Glu Glu Arg Leu Arg
 100 105 110
Ile Lys Leu Glu Pro Lys Gly Pro Gly Gly Pro Gln Gly Ser Trp Lys
 115 120 125
Trp Gly Gly Cys Ser Asp Asn Val Glu Phe Gly Ile Arg Phe Ser Arg
130 135 140
Glu Phe Val Asp Ala Arg Glu Arg Glu Lys Leu Met Thr Lys Ser Arg
145 150 155 160
Asp Arg Asp Ala Arg Ser Leu Met Asn Leu His Asn Asn Glu Ala Gly
 165 170 175
Arg Lys Ala Val Lys Ser His Met Arg Arg Glu Cys Lys Cys His Gly
 180 185 190
Val Ser Gly Ser Cys Ser Leu Lys Thr Cys Trp Leu Ser Leu Pro Asp
 195 200 205
Phe Arg Glu Val Gly Asp Leu Leu Lys Glu Lys Tyr Asp Gly Ala Ile
210 215 220
Glu Val Glu Val Asn Lys Arg Gly Lys Gly Gln Arg Ser Leu Ser Ser
225 230 235 240

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Arg	Lys	Gln	Ala	Ser 245	Ala	Leu	Glu	Ala	Ala 250	Asn	Glu	Arg	Phe	Lys 255	Lys
Pro	Thr	Arg	Asn 260	Gln	Tyr	Thr	Asp	Leu 265	Val	Tyr	Leu	Glu	Lys 270	Ser	Pro
Asp	Tyr	Cys 275	Glu	Arg	Asp	Arg	Glu 280	Thr	Gly	Ser	Leu	Gly 285	Thr	Gln	Gly
Arg	Val 290	Cys	Asn	Lys	Thr	Ser 295	Lys	Gly	Leu	Gln	Trp 300	Arg	Asp	Gly	Cys
Glu 305	Leu	Leu	Cys	Cys	Gly 310	Arg	Gly	Tyr	Asn	Thr 315	Glu	Gln	Lys	Val	Glu 320
Arg	Thr	Glu	Lys	Cys 325	Asn	Cys	Lys	Phe	His 330	Asn	Gly	Trp	Cys	Cys 335	Tyr
Val	Lys	Cys	Glu 340	Glu	Cys	Thr	Glu	Val 345	Val	Glu	Val	His	Thr 350	Cys	Lys

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<210> 72
<211> 1216
<212> PRT
<213> Rattus norvegicus
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<400>	72															
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Ala	Gln	Ser	Leu 20	Gly	Lys	Thr	Trp	Val 25	Pro	Asp	His	Cys	Arg 30	Ser	Pro	
Thr	Glu	Ala 35	Thr	Cys	Asn	Phe	Val 40	Cys	Asp	Cys	Gly	Asp 45	Cys	Ser	Asp	
Glu	Ala 50	Gln	Cys	Gly	Phe	His 55	Gly	Ala	Ser	Thr	Thr 60	Pro	Asn	Thr	Pro	
Phe 65	Thr	Cys	Asn	Phe	Glu 70	Gln	Asp	Pro	Cys	Gly 75	Trp	Gln	Asp	Ile	Ser 80	
Thr	Ser	Gly	Tyr	Arg 85	Trp	Leu	Arg	Asp	Arg 90	Ala	Gly	Ala	Gly	Leu 95	Asp	
Ser	Ser	Gly	Pro 100	His	Ser	Asp	His	Thr 105	Arg	Gly	Thr	Asp	Leu 110	Gly	Trp	
Tyr	Met	Ala 115	Val	Gly	Thr	His	Ser 120	Gly	Lys	Glu	Pro	Ser 125	Thr	Arg	Thr	
Leu	Arg 130	Ser	Pro	Val	Met	Arg 135	Glu	Ala	Ala	Pro	Thr 140	Cys	Glu	Leu	Arg	
Leu 145	Trp	Tyr	His	Thr	Asp 150	Ser	Arg	Asp	Val	Ala 155	Glu	Leu	Arg	Leu	Asp 160	
Leu	Thr	His	Gly	Met	Glu	Thr	Leu	Thr	Leu	Trp	Gln	Ser	Ser	Gly	Pro	

165		170		175
Trp Gly Pro Trp 180	Pro Gly Arg Glu Leu 185	Ala Val Asn Thr Gly 190	Arg Ile	
Gln Gly Asp 195	Phe Lys Val Thr Phe 200	Ser Ala Thr Arg Asn 205	Ala Thr His	
Arg Gly Ala Val 210	Ala Leu Asp 215	Asp Met Glu Phe Trp 220	Asp Cys Gly Leu	
Pro Ile Pro Gln Ala 225	Arg Cys Pro Leu Gly 230	His His His Cys Gln Asn 240		
Lys Ala Cys Val 245	Glu Pro His Gln Leu Cys 250	Asp Gly Glu Asp Asn Cys 255		
Gly Asp Ser 260	Ser Asp Glu Asp Pro Leu 265	Ile Cys Ser His His Met Ala 270		
Thr Asp Phe 275	Glu Thr Gly Leu Gly 280	Pro Trp Thr Gln Leu Glu Gly Trp 285		
Thr Arg Asn Phe 290	Ser Ala Gly 295	Ser Met Val Ser Pro Ala Trp Pro His 300		
Arg Asp His Ser Arg 305	Asn Ser Ala Tyr Gly 310	Phe Phe Leu Val Ser Val 320		
Ala Lys Pro Gly Thr 325	Thr Ala Val Leu Tyr 330	Ser Pro Glu Phe Gln Gly 335		
Ser Val Ser 340	Tyr Asn Cys Ser Phe Thr 345	Phe Tyr Tyr Tyr Leu His Gly 350		
Ser Glu Ala 355	Asn Gln Phe Gln Leu Phe 360	Val Gln Ala Gln Gly Leu Asn 365		
Thr Thr Gln Pro Pro 370	Val Leu Leu Arg Ser Arg 375	His Gly Glu Leu Gly 380		
Thr Ala Trp Val Arg 385	Asp Arg Val Asn Ile Gln 390	Ser Ala His Pro Phe 400		
Arg Ile Leu Leu Ala 405	Gly Glu Thr Gly Pro 410	Gly Gly Phe Val Gly Leu 415		
Asp Asp Leu Ile 420	Met Ser Asn His Cys 425	Ile Leu Val Pro Gly Met Ser 430		
Thr Leu Gln 435	Ser Ser Leu Ser Gly 440	Pro Val Pro Leu Ala Leu Tyr Pro 445		
Gln Thr Ser Ile Lys Arg 450	Thr Cys Asp Ala Gly 455	His Leu Ser Cys Asp 460		
Glu Leu Cys Val Pro 465	Pro Glu Gln Leu Cys 470	Asp Phe Gln Gln His Cys 475		
Ala Glu Gly Glu Asp 485	Glu Glu Lys Cys Gly 490	Thr Thr Asp Phe Glu Ser 495		
Ala Ser Ala Gly Gly Trp Glu Asp Ile Ser Ile Gly Lys Leu Gln Trp				

500					505					510					
Gln	Arg	Ala	Glu	Ala	Gln	Glu	Ser	Gly	Lys	Pro	Ala	Arg	Asp	Thr	Asn
		515					520					525			
Arg	Asn	Ala	Pro	Gly	His	Phe	Leu	Ser	Leu	Arg	Lys	Ala	Trp	Gly	Gln
	530					535					540				
Leu	Arg	Ser	Glu	Ala	Arg	Ala	Leu	Thr	Pro	Thr	Leu	Gly	Pro	Ser	Gly
545					550					555					560
Pro	His	Cys	Glu	Leu	His	Met	Thr	Tyr	Tyr	Phe	His	Ser	His	Pro	Gln
				565					570					575	
Gly	Phe	Leu	Ala	Leu	Ala	Val	Val	Glu	Asn	Gly	Phe	Arg	Glu	Leu	Leu
			580					585					590		
Trp	Gln	Ala	Pro	Ser	Ser	Ser	Ser	Gly	Gly	Trp	Thr	Leu	Gln	Lys	Ile
		595					600					605			
Leu	Leu	Gly	Ala	Arg	Arg	Trp	Pro	Phe	Gln	Leu	Glu	Phe	Val	Ser	Leu
	610					615					620				
Val	Asp	Leu	Asp	Gly	Pro	Gly	Gln	Gln	Gly	Ala	Gly	Val	Asp	Asn	Val
625					630					635					640
Thr	Leu	Arg	Asp	Cys	Asn	Pro	Met	Val	Thr	Thr	Glu	Ser	Asp	Gln	Glu
				645					650					655	
Val	Ser	Cys	Asn	Phe	Glu	Arg	Asp	Ser	Cys	Ser	Trp	His	Thr	Gly	His
			660					665					670		
Leu	Thr	Asp	Ala	His	Trp	His	Arg	Val	Lys	Ser	His	Gly	Ser	Gln	Tyr
		675					680					685			
Asp	His	Thr	Thr	Gly	Gln	Gly	Phe	Phe	Met	Phe	Leu	Asp	Pro	Met	Asp
	690					695					700				
Pro	Pro	Ala	Arg	Gly	Gln	Gly	Ala	Leu	Leu	Leu	Thr	Arg	Pro	Gln	Val
705					710					715					720
Pro	Val	Val	Pro	Lys	Glu	Cys	Leu	Ser	Phe	Trp	Tyr	His	Leu	His	Gly
				725					730					735	
Pro	Gln	Ile	Gly	Thr	Leu	Cys	Leu	Ala	Met	Arg	Arg	Glu	Gly	Glu	Glu
			740					745					750		
Asp	Thr	Leu	Leu	Trp	Ser	Arg	Ser	Gly	Thr	His	Gly	Asn	Arg	Trp	His
		755					760					765			
Gln	Ala	Trp	Val	Thr	Leu	His	His	Gln	Leu	Gln	Pro	Ser	Thr	Lys	Tyr
	770					775					780				
Gln	Leu	Leu	Phe	Glu	Gly	Leu	Arg	Asp	Gly	Tyr	His	Gly	Thr	Met	Gly
785					790					795					800
Leu	Asp	Asp	Met	Ala	Val	Arg	Pro	Gly	Pro	Cys	Trp	Ala	Ala	Lys	Arg
				805					810					815	
Cys	Ser	Phe	Glu	Asp	Ser	Asp	Cys	Gly	Phe	Ser	Pro	Gly	Asp	Trp	Gly
			820					825					830		
Leu	Trp	Thr	Arg	Gln	Asn	Asn	Ala	Ser	Gly	Leu	Gly	Pro	Trp	Gly	Pro

835 840 845
 Trp Ile Asp His Thr Thr Gly Thr Ala Gln Gly His Tyr Met Val Val
 850 855 860
 Asp Thr Ser Pro Asn Leu Leu Pro Lys Gly His Val Ala Ser Leu Thr
 865 870 875 880
 Ser Glu Glu His Pro Pro Leu Ser Arg Pro Ala Cys Leu Ser Phe Trp
 885 890 895
 Tyr His Leu Ser Phe His Asn Pro Gly Thr Leu Arg Val Phe Val Glu
 900 905 910
 Glu Ser Thr Arg Arg Gln Glu Leu Ser Ile Ser Gly His Gly Gly Phe
 915 920 925
 Ala Trp Arg Leu Gly Ser Val Asn Val Gln Ala Glu Gln Ala Trp Lys
 930 935 940
 Val Val Phe Glu Ala Met Ala Ser Gly Val Glu His Ser Tyr Met Ala
 945 950 955 960
 Leu Asp Asp Ile Ser Leu Gln Asp Gly Pro Cys Ala Gln Pro Gly Ser
 965 970 975
 Cys Asp Phe Glu Ser Gly Leu Cys Gly Trp Ser His Leu Pro Trp Pro
 980 985 990
 Gly Leu Gly Gly Tyr Ser Trp Asp Trp Ser Ser Gly Ala Thr Pro Ser
 995 1000 1005
 Arg Tyr Pro Arg Pro Ser Val Asp His Thr Val Gly Thr Glu Ala Gly
 1010 1015 1020
 His Phe Ala Phe Phe Glu Thr Ser Val Leu Gly Pro Gly Gly Gln Ala
 1025 1030 1035 1040
 Ala Trp Leu Gly Ser Glu Pro Leu Pro Ala Thr Ala Val Ser Cys Leu
 1045 1050 1055
 His Phe Trp Tyr Tyr Met Gly Phe Pro Ala His Phe Tyr Lys Gly Glu
 1060 1065 1070
 Leu Arg Val Leu Leu Ser Ser Thr Gln Gly Gln Leu Ala Val Trp His
 1075 1080 1085
 Arg Gly Gly His Leu Arg Asp Gln Trp Leu Gln Val Gln Ile Glu Val
 1090 1095 1100
 Ser Ser Ser Glu Glu Phe Gln Ile Val Phe Glu Ala Thr Leu Gly Gly
 1105 1110 1115 1120
 Gln Pro Ala Leu Gly Pro Ile Ala Leu Asp Asp Val Glu Tyr Leu Ala
 1125 1130 1135
 Gly Gln His Cys Lys Gln Pro Thr Pro Ser Gln Gly Arg Val Ala Ala
 1140 1145 1150
 Pro Val Ser Val Pro Val Ala Val Gly Gly Ala Leu Leu Leu Phe Leu
 1155 1160 1165
 Leu Leu Leu Gly Leu Gly Gly Trp His Trp Leu Gln Lys Gln His Leu

1170 1175
Pro Cys Gln Ser Thr Asp Ala Ala Ala Ser Gly Phe Asp Asn Ile Leu
1185 1190 1195 1200
Phe Asn Ala Asp Gln Val Thr Leu Pro Glu Ser Ile Thr Ser Asn Pro
1205 1210 1215

<210> 73
<211> 688
<212> PRT
<213> Xenopus laevis

<400> 73
Met Met Leu Ser His Trp Val Leu Leu Leu Ser Leu Gly Ala Val Trp
1 5 10 15
Leu Ala Glu Gly Gly Glu Ile Ser Pro Gly Ser Cys Thr Phe Glu Asn
20 25 30
Ser Thr Cys Ala Tyr Thr Ser Ala Phe Pro Phe Leu Gln Trp Thr Val
35 40 45
Asn Ile Glu Gly His Tyr Val Ser Val Asp Ser Ser Asn Gly Leu Arg
50 55 60
Gly Gln Lys Ala Val Leu Ile Ser Pro Asp Leu His Leu Ala Glu Trp
65 70 75 80
Ser Cys Leu Arg Leu Val Tyr Gln Ile Ala Gly Ser Glu Ser Ser Pro
85 90 95
Ser Pro Ser Ser Leu Asn Val Phe Val Arg Pro Glu Gly Glu Ser Phe
100 105 110
Asp Tyr Leu Leu Trp Ser Ala Glu Glu His Ser Asp Ser Trp Leu Ile
115 120 125
Ser Ser Ile Asp Leu Lys Asn Thr Thr Lys Arg Phe Lys Ile Ile Leu
130 135 140
Glu Gly Val Leu Gly Glu Asn Thr Met Ser Ser Ile Ala Ile Phe Glu
145 150 155 160
Val Lys Met Thr Thr Gly Tyr Cys Ile Glu Cys Asp Phe Glu Glu Asn
165 170 175
His Leu Cys Gly Tyr Met Asn Ser Trp Asn Pro Asn Val Asn Trp Phe
180 185 190
Val Gly Gly Gly Asn Val Lys Asn Ser His Ser Ile Leu Pro Arg Asp
195 200 205
His Thr Leu Asn Asn Glu Leu Gly His Tyr Met Tyr Val Asp Ser Val
210 215 220
Tyr Val Lys His Phe Gln Glu Val Ala Gln Leu Val Ser Pro Leu Ile
225 230 235 240

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Ile	Thr	Pro	Ile	Ser 245	Gly	Cys	Leu	Ser	Phe 250	Tyr	Tyr	Gln	Leu	Gln 255	Arg
Glu	Thr	Ser	Asn 260	Ile	Phe	Leu	Val	His 265	Thr	Arg	Asp	Leu	His 270	Gly	Ser
Tyr	Asp	Glu 275	Ile	Trp	Lys	Met	Gly 280	Ala	Val	Arg	Gln	Gly 285	Glu	Trp	Asn
Leu	Ala 290	Glu	Val	Asp	Leu	Asn 295	Ala	His	Val	Pro	Leu 300	Glu	Val	Ile	Phe
Glu 305	Val	Ala	Phe	Asn	Gly 310	Ile	Gln	Ala	Gly	Tyr 315	Val	Ala	Leu	Asp	Asp 320
Ile	Leu	Phe	Ser	Pro 325	Val	Ser	Cys	Ser	Gly 330	Gln	Glu	Gly	Met	Phe 335	Phe
Asp	Ala	Arg	Glu 340	Ala	Gly	Cys	Asp	Phe 345	Glu	Glu	Gly	Met	Cys 350	Gln	Phe
His	Gln	Asp 355	Asp	Asn	Asn	Gly	Ser 360	Gly	Trp	Ser	Arg	Val 365	Lys	Val	Lys
Pro	Asn 370	Ala	Tyr	Gln	Met	Gly 375	Asp	His	Thr	Thr	Gly 380	Leu	Gly	Tyr	Phe
Met 385	Ile	Ala	Asn	Thr	Arg 390	Phe	Thr	Gly	Gln	Pro 395	Ala	Tyr	Phe	Gly	Arg 400
Leu	Tyr	Gly	Pro	Ser 405	Leu	Pro	Gly	Asn	Ile 410	Gln	Tyr	Cys	Ile	Arg 415	Phe
Phe	Tyr	Ser	Leu 420	Tyr	Gly	Phe	Tyr	Lys 425	Thr	Ile	Asp	Ser	Leu 430	Ala	Val
Tyr	Ile	Phe 435	Glu	Glu	Asn	His	Val 440	Val	Gln	Glu	Lys	Ile 445	Trp	Ser	Ala
His	Glu 450	Thr	Pro	Lys	Gly	Val 455	Trp	Leu	Gln	Ala	Glu 460	Ile	Ser	Ile	His
Lys 465	Pro	Met	Pro	Phe	Lys 470	Val	Val	Phe	Val	Ser 475	Trp	Cys	Lys	Ser	Leu 480
Trp	Asp	Cys	Gly	Ile 485	Ala	Ala	Leu	Asp	Asp 490	Ile	Ser	Val	Ser	Ile 495	Gly
Ser	Cys	Lys	Ile 500	Ser	Asp	Arg	Ile	Pro 505	Pro	Leu	Pro	Gly	Lys 510	Cys	Thr
Phe	Glu	Lys 515	Asn	Asp	Cys	Gly	Phe 520	Gly	Ala	Gly	Met	Ala 525	Lys	Glu	Gly
Tyr	Leu 530	Ala	Gln	Asn	Thr	Arg 535	Glu	Asp	Pro	Thr	Phe 540	Tyr	Thr	Gly	Pro
Asn 545	Gly	Asp	His	Thr	Ser 550	Gly	Val	Gly	Tyr	Tyr 555	Met	Tyr	Ile	Glu	Ala 560
Thr	Asn	Met	Val	Phe 565	Gly	Gln	Lys	Ala	Lys 570	Leu	Ile	Ser	Arg	Pro 575	Leu

Arg Ala Val Ala Gly Lys Gln Cys Leu Thr Phe Tyr Tyr His Met Tyr
 580 585 590
 Gly Ala Gly Thr Gly Leu Leu Asn Val Tyr Leu Thr Lys Glu Gly Asp
 595 600 605
 Ile Asn Lys Asp Thr Leu Leu Trp Thr Arg Lys Gly Glu Gln Ser Ile
 610 615 620
 Thr Trp Leu Lys Ala Gln Met Glu Tyr Glu Ser Glu Gln Gln His Lys
 625 630 635 640
 Ile Val Phe Glu Ala Val Arg Gly Ile Ser Ile Arg Ser Asp Ile Ala
 645 650 655
 Ile Asp Asp Ile Leu Phe Gln Asn Gly Pro Cys Asn Asp Ser Ser Asp
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 Pro Leu Gln Ser Ser Gly Tyr Ser Asp Asn Phe Asn Asn Ile Glu Phe
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 <211> 5376
 <212> PRT
 <213> Mus musculus

<400> 74
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 Gly Gln Glu Gln Val Pro Ala Trp Arg Pro Asn Ser Pro Asp Leu Gly
 20 25 30
 Pro Met Val His Thr Ser Arg Glu Asp Ser Ile Leu Ser Lys Cys Asp
 35 40 45
 Phe Glu Asp Asn Ser Arg Pro Phe Cys Asp Trp Ser Gln Met Ser Ala
 50 55 60
 Asp Asp Gly Asp Trp Ile Arg Thr Thr Gly Pro Ser Leu Thr Gly Thr
 65 70 75 80
 Ser Gly Pro Pro Gly Gly Tyr Pro Asn Gly Glu Gly Tyr Tyr Leu His
 85 90 95
 Met Asp Pro Lys Thr Phe Pro Gln Gly Gly Val Ala Arg Leu Arg Ser
 100 105 110
 Pro Asp Ile Trp Glu Gln Gly Pro Leu Cys Val His Phe Ala Phe His
 115 120 125
 Met Phe Gly Leu Ser Trp Gly Ala Gln Leu Arg Leu Leu Leu Arg
 130 135 140
 Gly Arg Lys His Leu Arg Pro Tyr Val Leu Trp Lys His Val Asn Thr
 145 150 155 160
 Gln Ser Pro Ser Trp Met Pro Thr Thr Val Thr Val Pro Ala Asp His
 165 170 175

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Asp Ile Pro Ser Trp Leu Met Phe Glu Gly Met Arg Gly Asn Thr Ala
 180 185 190
 Tyr Leu Asp Ile Ser Leu Asp Gly Leu Ser Ile Gln Arg Gly Thr Cys
 195 200 205
 Asn Gln Val Cys Met Ser Gln Met Cys Thr Phe Asp Thr Leu Asn Asp
 210 215 220
 Leu Cys Gly Trp Ser Trp Val Pro Thr Ala Thr Gly Ala Lys Trp Thr
 225 230 235 240
 Gln Lys Lys Gly Pro Thr Gly Lys Gln Gly Val Gly Pro Ala Glu Asp
 245 250 255
 Phe Ser Asn Pro Gly Asn Gly Tyr Tyr Met Leu Leu Asp Ser Thr Asn
 260 265 270
 Ala Arg Pro Gly Gln Lys Ala Val Leu Leu Ser Pro Leu Ser His Ser
 275 280 285
 Arg Gly Cys Met Thr Leu Ser Phe His Tyr Ile Met His Gly Gln Gly
 290 295 300
 His Glu Glu Gly Leu Phe Val Tyr Ala Thr Phe Leu Gly Asn Ile Arg
 305 310 315 320
 Lys Tyr Thr Leu Phe Ser Gly His Pro Gly Pro Asp Trp Gln Ala Val
 325 330 335
 Ser Val Asn Tyr Thr Gly Gln Gly Gln Ile Gln Phe Met Val Val Gly
 340 345 350
 Met Phe Gly Asn Ile Pro Glu Pro Ala Ile Ala Val Asp Ala Ile Ser
 355 360 365
 Ile Ala Pro Cys Gly Glu Ser Phe Pro Gln Cys Asp Phe Glu Asp Arg
 370 375 380
 Val His Pro Phe Cys Asp Trp Asn Gln Val Tyr Gly Asp Met Gly His
 385 390 395 400
 Trp Ser Trp Gly Ser Lys Ser Val Pro Thr Leu Ile Ala Gly Ser Pro
 405 410 415
 Arg Glu Phe Pro Tyr Gly Gly Glu His Tyr Ile Phe Phe Asp Ser Val
 420 425 430
 Lys Leu Ser Gln Glu Gly Gln Ser Ala Arg Leu Val Ser Pro Pro Phe
 435 440 445
 Cys Ala Pro Gly Gly Ile Cys Val Glu Phe Ala Tyr His Met Tyr Gly
 450 455 460
 Leu Gly Lys Gly Thr Thr Leu Lys Leu Leu Leu Gly Ser Pro Ala Gly
 465 470 475 480
 Ser Ser Pro Ile Pro Leu Trp Asn Arg Val Gly Ser Gln Ser Ser Gly
 485 490 495
 Trp Met Asn Ser Ser Val Thr Ile Pro Lys Gly Tyr Gln Gln Pro Met
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Gln Leu Phe Ile Glu Ala Thr Arg Gly Thr Ser Thr Ala Phe Val Val
 515 520 525
 Ala Leu Asn Phe Ile Leu Ile Ser His Gly Pro Cys Arg Val Leu Leu
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 Gln Thr Glu Ile Pro Ser Ser Pro Leu Leu Pro Pro Thr Gly Pro Ser
 545 550 555 560
 Glu Ser Thr Val Pro Thr Leu Pro Met Glu Gln Pro Thr Ser Pro Thr
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 Lys Ala Thr Thr Val Thr Ile Glu Ile Pro Thr Thr Pro Thr Glu Glu
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 Ala Thr Ile Pro Thr Glu Thr Thr Thr Val Pro Thr Glu Val Ile Asn
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 Val Ser Pro Lys Glu Thr Ser Ile Pro Pro Glu Val Thr Ile Pro Thr
 610 615 620
 Glu Val Ile Thr Val Ser Pro Glu Glu Ile Ile Ser Pro Thr Glu Val
 625 630 635 640
 Thr Pro Val Pro Thr Asp Val Thr Ala Ala Tyr Val Glu Ala Thr Asn
 645 650 655
 Ala Ser Pro Glu Glu Thr Ser Val Pro Pro Glu Val Thr Ile Leu Thr
 660 665 670
 Glu Val Thr Thr Val Ser Pro Glu Glu Thr Thr Val Pro Thr Glu Val
 675 680 685
 Pro Ile Val Leu Ile Glu Ala Thr Ala Phe Pro Thr Gly Glu Thr Thr
 690 695 700
 Leu Tyr Thr Glu Val Pro Thr Val Pro Thr Glu Val Thr Gly Val His
 705 710 715 720
 Thr Glu Val Thr Asn Val Ser Pro Glu Glu Thr Ser Val Pro Thr Glu
 725 730 735
 Glu Thr Ile Ser Thr Glu Val Thr Thr Val Ser Pro Glu Glu Thr Thr
 740 745 750
 Val Pro Thr Glu Val Pro Ile Val Leu Ile Glu Ala Thr Ala Ser Pro
 755 760 765
 Thr Gly Glu Ile Thr Leu Tyr Thr Glu Val Pro Thr Val Pro Thr Glu
 770 775 780
 Val Thr Gly Val His Thr Glu Val Thr Asn Val Ser Pro Glu Glu Thr
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 Ser Val Pro Thr Glu Glu Thr Ile Ser Thr Glu Val Thr Thr Val Ser
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 Pro Glu Glu Thr Thr Leu Pro Thr Glu Val Pro Thr Val Ser Thr Glu
 820 825 830
 Val Thr Asn Val Ser Pro Glu Glu Thr Ser Val Pro Pro Glu Glu Thr
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Ile Leu Thr Thr Leu Tyr Thr Glu Val Pro Thr Val Pro Thr Glu Val
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 Thr Gly Val His Thr Glu Val Thr Asn Val Ser Pro Glu Glu Thr Ser
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 Val Pro Thr Glu Glu Thr Ile Ser Thr Glu Val Thr Thr Val Ser Pro
 885 890 895
 Glu Glu Thr Thr Leu Pro Thr Glu Val Pro Thr Val Ser Thr Glu Val
 900 905 910
 Thr Asn Val Ser Pro Glu Glu Thr Ser Val Pro Pro Glu Glu Thr Ile
 915 920 925
 Leu Thr Glu Ile Thr Thr Val Ser Pro Glu Glu Thr Val Phe Pro Ile
 930 935 940
 Glu Gly Thr Thr Leu Pro Thr Glu Val Leu Thr Val Pro Ile Glu Val
 945 950 955 960
 Thr Thr Phe Pro Thr Gly Glu Thr Thr Val Pro Thr Glu Val Pro Thr
 965 970 975
 Val Ser Thr Glu Met Thr Gly Val His Thr Glu Val Thr Thr Val Phe
 980 985 990
 Pro Glu Glu Thr Ser Ile Pro Thr Glu Val Ala Thr Val Leu Pro Ala
 995 1000 1005
 Ser Ile Pro Pro Glu Glu Thr Thr Thr Pro Thr Glu Val Thr Thr Thr
 1010 1015 1020
 Pro Pro Glu Glu Thr Thr Ile Pro Ala Glu Val Thr Thr Val Pro Pro
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 Ala Ser Ile Pro Pro Glu Glu Thr Ala Ser Leu Thr Glu Val Thr Thr
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 Thr Pro Pro Glu Glu Thr Thr Thr Pro Thr Glu Val Thr Thr Val Pro
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 Pro Glu Lys Thr Thr Ile Pro Thr Glu Val Thr Thr Val Pro Pro Ala
 1075 1080 1085
 Ser Ile Phe Pro Glu Glu Thr Thr Val Pro Pro Glu Glu Thr Thr Ile
 1090 1095 1100
 Ala Ser Glu Glu Thr Thr Val Ser Thr Gln Glu Thr Thr Leu Leu Thr
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 Glu Gln Ser Ala Val Thr Gln Thr Ser Ile Ala Cys Arg Pro Pro Cys
 1125 1130 1135
 Pro Ser Pro Pro Leu Met Pro Ile Gly Pro Leu Leu Ser Lys Pro Pro
 1140 1145 1150
 Gly Val Ser Met Phe Ser Leu Ala Pro Thr Thr Gly Val Ser Thr Thr
 1155 1160 1165
 Glu Ser Cys Pro Pro Asn Ala His Ile Glu Leu Cys Ala Cys Pro Ala
 1170 1175 1180

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Ser Cys Glu Ser Pro Lys Pro Ser Cys Gln Pro Pro Cys Ile Pro Gly
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 Cys Val Cys Asn Pro Gly Phe Leu Phe Ser Asn Asn Gln Cys Ile Asn
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 Glu Ser Ser Cys Asn Cys Pro Tyr Asn Asn Lys His Tyr Lys Pro Gly
 1220 1225 1230
 Glu Glu Trp Phe Thr Pro Asn Cys Thr Glu Arg Cys Arg Cys Leu Pro
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 Gly Ser Leu Met Glu Cys Gln Ile Ser Gln Cys Gly Thr His Thr Val
 1250 1255 1260
 Cys Gln Leu Lys Ser Asp Gln Tyr Gln Cys Glu Pro Tyr Gly Lys Ala
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 Thr Cys Leu Val Tyr Gly Asp Leu His Phe Val Thr Phe Asp Glu Arg
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 His Ile Gly Phe Thr Gly Thr Cys Thr Tyr Ile Leu Thr Gln Thr Cys
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 Ser Asn Ser Thr Asp His Phe Phe Arg Ile Thr Ala Asn Thr Glu Glu
 1315 1320 1325
 Arg Gly Val Glu Gly Val Ser Cys Leu Asp Lys Val Val Ile Ser Leu
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 Pro Glu Thr Thr Val Thr Met Ile Ser Gly Arg His Thr Leu Ile Gly
 1345 1350 1355 1360
 Asp Gln Glu Val Thr Leu Pro Ala Ile Leu Ser Asp Asp Thr Tyr Val
 1365 1370 1375
 Gly Leu Ser Gly Arg Phe Val Glu Leu Arg Thr Thr Phe Gly Leu Arg
 1380 1385 1390
 Val Arg Trp Asp Gly Asp Gln Gln Leu Phe Val Thr Val Ser Ser Thr
 1395 1400 1405
 Phe Ser Gly Lys Leu Cys Gly Phe Cys Gly Asn Tyr Asp Gly Asp Ser
 1410 1415 1420
 Ser Asn Asp Asn Leu Lys Ser Asp Gly Met Met Thr His Asp Glu Glu
 1425 1430 1435 1440
 Glu Leu Arg Leu Ser Trp Gln Val Glu Glu Asp Glu Asp Lys Asp Trp
 1445 1450 1455
 Val Ser Ser Arg Cys Gln Lys Lys Lys Asn Pro Pro Ser Cys Asp Ala
 1460 1465 1470
 Ala Leu Gly Ser Thr Met Ser Gly Pro Lys Leu Cys Gly Gln Leu Val
 1475 1480 1485
 Asn Pro Ser Gly Pro Phe Glu Ala Cys Leu Leu His Leu Lys Ala Ser
 1490 1495 1500
 Ser Phe Leu Asp Asn Cys Val Thr Asp Met Cys Ser Phe Gln Gly Leu
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Gln Gln Lys Leu Cys Ala Arg Met Ser Ala Met Thr Ala Thr Cys Gln
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 Asp Ala Gly Tyr Pro Val Lys Pro Trp Arg Glu Pro Gln Phe Cys Pro
 1540 1545 1550
 Leu Val Cys Pro Lys Asn Ser Arg Tyr Ser Leu Cys Ala Lys Pro Cys
 1555 1560 1565
 Pro Glu Thr Cys His Pro Ile Ser Thr Thr Gln His Cys Ser Asp Lys
 1570 1575 1580
 Cys Val Glu Gly Cys Glu Cys Asp Pro Gly Phe Ile Leu Ser Gly Ser
 1585 1590 1595 1600
 Glu Cys Val Pro Ser Ser Gln Cys Gly Cys Thr Ser Phe Gln Gly Arg
 1605 1610 1615
 Tyr Phe Lys Leu Gln Glu Gln Trp Phe Asn Pro Asp Cys Lys Glu Ile
 1620 1625 1630
 Cys Thr Cys Glu Ser His Asn His Ile Leu Cys Lys Pro Trp Lys Cys
 1635 1640 1645
 Lys Ala Gln Glu Ala Cys Ser Tyr Lys Asn Gly Val Leu Gly Cys His
 1650 1655 1660
 Ala Gln Gly Ala Ala Thr Cys Met Val Ser Gly Asp Pro His Tyr Leu
 1665 1670 1675 1680
 Thr Phe Asp Gly Ala Leu His His Phe Met Gly Thr Cys Thr Tyr Val
 1685 1690 1695
 Leu Thr Gln Pro Cys Trp Ser Lys Ser Gln Glu Asn Asn Phe Val Val
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 Ser Ala Thr Asn Glu Ile His Asp Gly Asn Leu Glu Val Ser Tyr Val
 1715 1720 1725
 Lys Ala Val His Val Gln Val Phe Asp Leu Lys Ile Ser Met Phe Lys
 1730 1735 1740
 Gly Gln Lys Val Val Leu Asn Asn Gln Arg Val Val Leu Pro Val Trp
 1745 1750 1755 1760
 Pro Ser Gln Gly Arg Val Thr Ile Arg Leu Ser Gly Ile Phe Val Leu
 1765 1770 1775
 Leu Tyr Thr Asn Phe Gly Leu Gln Val Arg Tyr Asp Gly Arg His Leu
 1780 1785 1790
 Val Glu Val Thr Val Pro Ser Ser Tyr Thr Gly Ser Leu Cys Gly Leu
 1795 1800 1805
 Cys Gly Asn Tyr Asn Asn Asn Ser Met Asp Asp Asn Leu Arg Ala Asp
 1810 1815 1820
 Met Lys Pro Ala Gly Asn Ser Leu Leu Leu Gly Ala Ala Trp Lys Ile
 1825 1830 1835 1840
 Leu Glu Ala Ser Asp Pro Gly Cys Phe Leu Ala Gly Gly Lys Pro Ser
 1845 1850 1855

Arg Cys Ala Asp Ser Asp Met Asp Asp Val Trp Thr Lys Lys Cys Ala
 1860 1865 1870
 Ile Leu Met Asn Pro Leu Gly Pro Phe Ser Asn Cys His Glu Ala Val
 1875 1880 1885
 Pro Pro Gln Ala Ser Phe Ser Ser Cys Val Tyr Gly Gln Cys Glu Thr
 1890 1895 1900
 Asn Gly Asp Asn Leu Thr Phe Cys His Ser Leu Gln Ala Tyr Ala Ser
 1905 1910 1915 1920
 Leu Cys Ala Gln Ala Gly Gln Val Thr Thr Trp Arg Asn Ser Thr Phe
 1925 1930 1935
 Cys Pro Met Arg Cys Pro Pro Arg Ser Ser Tyr Asn Pro Cys Ala Asn
 1940 1945 1950
 Ser Cys Pro Ala Thr Cys Leu Thr Leu Ser Thr Pro Arg Asp Cys Pro
 1955 1960 1965
 Thr Leu Pro Cys Val Glu Gly Cys Glu Cys Gln Ser Gly His Ile Leu
 1970 1975 1980
 Ser Gly Thr Thr Cys Val Pro Leu Arg Gln Cys Gly Cys Ser Asp Gln
 1985 1990 1995 2000
 Asp Gly Ser Tyr His Leu Leu Gly Glu Ser Trp Tyr Thr Glu Lys Thr
 2005 2010 2015
 Cys Thr Thr Leu Cys Thr Cys Ser Ala His Ser Asn Ile Thr Cys Ser
 2020 2025 2030
 Pro Thr Ala Cys Lys Ala Asn His Val Cys Leu Arg Gln Glu Gly Leu
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 Leu Arg Cys Ala Ala Glu Met Gly Glu Cys Arg Ile Ser Glu Asp Ser
 2050 2055 2060
 Gln Ile Val Ser Phe Asp Asp His Ser His Pro Ile Gln Asp Thr Cys
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 Thr Tyr Ile Leu Val Lys Val Cys His Pro Asn Thr Asn Met Pro Phe
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 Phe Met Ile Ser Ala Lys Thr Asp Ile Asn Thr Asn Gly Lys Asn Lys
 2100 2105 2110
 Thr Phe Gly Val Tyr Gln Leu Tyr Ile Asp Ile Phe Asn Phe His Ile
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 Thr Leu Gln Lys Asp His Leu Val Leu Ile Ser Leu Ile Asn Asp Ser
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 Ile Val Thr Leu Pro Thr Thr Thr His Ile Pro Gly Val Ser Val Met
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 Thr Glu Asp Val Tyr Thr Ile Val Thr Ile Lys Asp Glu Ile Gln Val
 2165 2170 2175
 Lys Phe Glu Ser Asn Asn Phe Leu Asp Val Lys Ile Pro Ala Ser Ser
 2180 2185 2190

Asn Gly Lys Val Cys Gly Val Cys Gly Asn Phe Asn Gly Glu Glu Glu
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 Asp Glu Leu Met Thr Pro Ser Gly Glu Leu Ala Glu Asp Glu Gln Glu
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 Phe Met Asn Ser Trp Lys Asp Lys Ser Met Asp Pro Asn Cys Gln Lys
 2225 2230 2235 2240
 Ile Glu Gly Gln Asn Leu Gln Val Glu Gln Gln Glu Ile Met Asn Gly
 2245 2250 2255
 Lys Cys Arg Pro Ile Asp Phe Glu Lys Ala Gln Ala Asn Cys Gln Thr
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 Lys Pro Phe Leu Leu Lys Cys Met Asn Ser Phe Cys Glu Phe Arg Glu
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 Leu Phe Arg Ala Leu Cys Asp Ser Leu Gln Ser Phe Glu Asp Ala Cys
 2305 2310 2315 2320
 Gln Asn Gln Gly Leu Lys Pro Pro Ile Trp Arg Asn Ser Ser Phe Cys
 2325 2330 2335
 Pro Leu Glu Cys Pro Ala His Ser His Tyr Thr Asn Cys Leu Pro Ser
 2340 2345 2350
 Cys Pro Pro Ser Cys Leu Asp Pro Asp Ser Arg Cys Glu Gly Ser Gly
 2355 2360 2365
 His Lys Val Pro Ala Thr Cys Arg Glu Gly Cys Ile Cys Gln Pro Asp
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 Tyr Val Leu Leu Asn Asp Lys Cys Val Leu Arg Ser His Cys Gly Cys
 2385 2390 2395 2400
 Lys Asp Ala Gln Gly Val Phe Ile Pro Ala Gly Lys Thr Trp Ile Ser
 2405 2410 2415
 Glu Asp Cys Thr Gln Ser Cys Thr Cys Met Lys Gly Ser Met Arg Cys
 2420 2425 2430
 Trp Asp Phe Gln Cys Pro Pro Gly Thr Tyr Cys Lys Asn Ser Asn Asp
 2435 2440 2445
 Gly Ser Ser Asn Cys Val Lys Ile Ser Leu Gln Cys Pro Ala His Ser
 2450 2455 2460
 Lys Phe Thr Asp Cys Leu Pro Pro Cys His Pro Ser Cys Ser Asp Pro
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 Asp Gly His Cys Glu Gly Ile Ser Thr Asn Ala His Ser Asn Cys Lys
 2485 2490 2495
 Glu Gly Cys Val Cys Gln Pro Gly Tyr Val Leu Arg Asn Asp Lys Cys
 2500 2505 2510
 Val Leu Arg Ile Glu Cys Gly Cys Gln His Thr Gln Gly Gly Phe Ile
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Pro Ala Gly Lys Asn Trp Thr Ser Arg Gly Cys Ser Gln Ser Cys Asp
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 2545 2550 2555 2560
 Thr Tyr Cys Gln Asp Ile Glu Asp Gly Thr Ser Asn Cys Ala Asn Ile
 2565 2570 2575
 Thr Leu Gln Cys Pro Ala His Ser Ser Phe Thr Asn Cys Leu Pro Pro
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 Cys Gln Pro Ser Cys Ser Asp Pro Glu Gly His Cys Gly Gly Ser Thr
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 Thr Lys Ala Pro Ser Ala Cys Gln Glu Gly Cys Val Cys Glu Pro Asp
 2610 2615 2620
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 2645 2650 2655
 Lys Gly Cys Thr Gln Thr Cys Ala Cys Val Thr Gly Thr Ile His Cys
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 Arg Asp Phe Gln Cys Pro Ser Gly Thr Tyr Cys Lys Asp Ile Lys Asp
 2675 2680 2685
 Asp Ala Ser Asn Cys Thr Glu Ile Ile Leu Gln Cys Pro Asp His Ser
 2690 2695 2700
 Leu Tyr Thr His Cys Leu Pro Ser Cys Leu Leu Ser Cys Ser Asp Pro
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 Asp Gly Leu Cys Arg Gly Thr Ser Pro Glu Ala Pro Ser Thr Cys Lys
 2725 2730 2735
 Glu Gly Cys Val Cys Asp Pro Asp Tyr Val Leu Ser Asn Asp Lys Cys
 2740 2745 2750
 Val Leu Arg Ile Glu Cys Gly Cys Lys Asp Ala Gln Gly Val Leu Ile
 2755 2760 2765
 Pro Ala Gly Lys Thr Trp Ile Asn Arg Gly Cys Thr Gln Ser Cys Ser
 2770 2775 2780
 Cys Met Gly Gly Ala Ile Gln Cys Gln Asn Phe Lys Cys Pro Ser Glu
 2785 2790 2795 2800
 Ala Tyr Cys Gln Asp Met Glu Asp Gly Asn Ser Asn Cys Thr Ser Ile
 2805 2810 2815
 Pro Leu Gln Cys Pro Ala His Ser His Tyr Thr Asn Cys Leu Pro Thr
 2820 2825 2830
 Cys Gln Pro Ser Cys Ser Asp Pro Asp Gly His Cys Glu Gly Ser Ser
 2835 2840 2845
 Thr Lys Ala Pro Ser Ala Cys Lys Glu Gly Cys Val Cys Glu Pro Asp
 2850 2855 2860

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Tyr Val Met Leu Asn Asn Lys Cys Val Pro Arg Ile Glu Cys Gly Cys
 2865 2870 2875 2880
 Lys Asp Thr Gln Gly Val Leu Ile Pro Ala Asp Lys Thr Trp Ile Asn
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 Arg Gly Cys Thr Gln Ser Cys Thr Cys Arg Gly Gly Ala Ile Gln Cys
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 Gln Lys Tyr His Cys Ser Ser Gly Thr Tyr Cys Lys Asp Met Glu Asp
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 Asp Ser Ser Ser Cys Ala Thr Ile Thr Leu Gln Cys Pro Ala His Ser
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 His Phe Thr Asn Cys Leu Pro Pro Cys Gln Pro Ser Cys Leu Asp Ser
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 Glu Gly His Cys Glu Gly Ser Thr Thr Lys Ala Pro Ser Ala Cys Gln
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 Val Pro Arg Ile Glu Cys Gly Cys Lys Asp Ala Gln Gly Val Leu Ile
 2995 3000 3005
 Pro Ala Asp Lys Thr Trp Ile Asn Arg Gly Cys Thr Gln Ser Cys Thr
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 Ser Leu Gln Cys Pro Ala Asn Ser Asn Phe Thr Ser Cys Leu Pro Ser
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 Cys Gln Pro Ser Cys Ser Asn Thr Asp Val His Cys Glu Gly Ser Ser
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 Pro Asn Thr Leu Ser Ser Cys Arg Glu Gly Cys Val Cys Gln Ser Gly
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 Tyr Val Leu His Asn Asp Lys Cys Ile Leu Arg Asn Gln Cys Gly Cys
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 Lys Asp Ala Gln Gly Ala Leu Ile Pro Glu Gly Lys Thr Trp Ile Thr
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 Ser Gly Cys Thr Gln Ser Cys Asn Cys Thr Gly Gly Ala Ile Gln Cys
 3140 3145 3150
 Gln Asn Phe Gln Cys Pro Leu Lys Thr Tyr Cys Lys Asp Leu Lys Asp
 3155 3160 3165
 Gly Ser Ser Asn Cys Thr Asn Ile Pro Leu Gln Cys Pro Ala His Ser
 3170 3175 3180
 Arg Tyr Thr Asn Cys Leu Pro Ser Cys Pro Pro Leu Cys Leu Asp Pro
 3185 3190 3195 3200

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Glu Gly Leu Cys Glu Gly Thr Ser Pro Lys Val Pro Ser Thr Cys Arg
 3205 3210 3215
 Glu Gly Cys Ile Cys Gln Pro Gly Tyr Leu Met His Lys Asn Lys Cys
 3220 3225 3230
 Val Leu Arg Ile Phe Cys Gly Cys Lys Asn Thr Gln Gly Ala Phe Ile
 3235 3240 3245
 Ser Ala Asp Lys Thr Trp Ile Ser Arg Gly Cys Thr Gln Ser Cys Thr
 3250 3255 3260
 Cys Pro Ala Gly Ala Ile His Cys Arg Asn Phe Lys Cys Pro Ser Gly
 3265 3270 3275 3280
 Thr Tyr Cys Lys Asn Gly Asp Asn Gly Ser Ser Asn Cys Thr Glu Ile
 3285 3290 3295
 Thr Leu Gln Cys Pro Thr Asn Ser Gln Phe Thr Asp Cys Leu Pro Ser
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 Cys Val Pro Ser Cys Ser Asn Arg Cys Glu Val Thr Ser Pro Ser Val
 3315 3320 3325
 Pro Ser Ser Cys Arg Glu Gly Cys Leu Cys Asn His Gly Phe Val Phe
 3330 3335 3340
 Ser Glu Asp Lys Cys Val Pro Arg Thr Gln Cys Gly Cys Lys Asp Ala
 3345 3350 3355 3360
 Arg Gly Ala Ile Ile Pro Ala Gly Lys Thr Trp Thr Ser Lys Gly Cys
 3365 3370 3375
 Thr Gln Ser Cys Ala Cys Val Glu Gly Asn Ile Gln Cys Gln Asn Phe
 3380 3385 3390
 Gln Cys Pro Pro Glu Thr Tyr Cys Lys Asp Asn Ser Glu Gly Ser Ser
 3395 3400 3405
 Thr Cys Thr Lys Ile Thr Leu Gln Cys Pro Ala His Thr Gln Tyr Thr
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 Ser Cys Leu Pro Ser Cys Leu Pro Ser Cys Leu Asp Pro Glu Gly Leu
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 Cys Lys Asp Ile Ser Pro Lys Val Pro Ser Thr Cys Lys Glu Gly Cys
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 Val Cys Gln Ser Gly Tyr Val Leu Asn Ser Asp Lys Cys Val Leu Arg
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 Lys Thr Trp Thr Ser Pro Gly Cys Thr Gln Ser Cys Ala Cys Met Gly
 3490 3495 3500
 Gly Ala Val Gln Cys Gln Ser Ser Gln Cys Pro Pro Gly Thr Tyr Cys
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 Lys Asp Asn Glu Asp Gly Asn Ser Asn Cys Ala Lys Ile Thr Leu Gln
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Cys Pro Ala His Ser Leu Phe Thr Asn Cys Leu Pro Pro Cys Leu Pro
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 Ser Cys Leu Asp Pro Asp Gly Leu Cys Lys Gly Ala Ser Pro Lys Val
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 Pro Ser Thr Cys Lys Glu Gly Cys Ile Cys Gln Ser Gly Tyr Val Leu
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 Ser Asn Asn Lys Cys Leu Leu Arg Asn Arg Cys Gly Cys Lys Asp Ala
 3585 3590 3595 3600
 His Gly Ala Leu Ile Pro Glu Asp Lys Thr Trp Val Ser Arg Gly Cys
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 Thr Gln Ser Cys Val Cys Thr Gly Gly Ser Ile Gln Cys Leu Ser Ser
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 Gln Cys Pro Pro Gly Ala Tyr Cys Lys Asp Asn Glu Asp Gly Ser Ser
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 Asn Cys Ala Arg Ile Pro Pro Gln Cys Pro Ala Asn Ser His Tyr Thr
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 Asp Cys Phe Pro Pro Cys Pro Pro Ser Cys Ser Asp Pro Glu Gly His
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 Cys Glu Ala Ser Gly Pro Arg Val Leu Ser Thr Cys Arg Glu Gly Cys
 3685 3690 3695
 Leu Cys Asn Pro Gly Phe Val Leu Asp Arg Asp Lys Cys Val Pro Arg
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 Val Glu Cys Gly Cys Lys Asp Ala Gln Gly Ala Leu Ile Pro Ser Gly
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 Lys Thr Trp Thr Ser Pro Gly Cys Thr Gln Ser Cys Ala Cys Met Gly
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 Gly Val Val Gln Cys Gln Ser Ser Gln Cys Pro Pro Gly Thr Tyr Cys
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 Lys Asp Asn Glu Asp Gly Asn Ser Asn Cys Ala Lys Ile Thr Leu Gln
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 Cys Pro Thr His Ser Asn Tyr Thr Asp Cys Leu Pro Phe Cys Leu Pro
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 Ser Cys Leu Asp Pro Ser Ala Leu Cys Gly Gly Thr Ser Pro Lys Gly
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 Pro Ser Thr Cys Lys Glu Gly Cys Val Cys Gln Pro Gly Tyr Val Leu
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 3825 3830 3835 3840
 Gln Gly Ala Val Ile Pro Ala Gly Lys Thr Trp Leu Ser Thr Gly Cys
 3845 3850 3855
 Ile Gln Ser Cys Ala Cys Val Glu Gly Thr Ile Gln Cys Gln Asn Phe
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Gln Cys Pro Pro Gly Thr Tyr Cys Asn His Asn Asn Asn Cys Ala Lys
 3875 3880 3885
 Ile Pro Leu Gln Cys Pro Ala His Ser His Phe Thr Ser Cys Leu Pro
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 Ser Cys Pro Pro Ser Cys Ala Asn Leu Asp Gly Ser Cys Glu Gln Thr
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 Gly Tyr Phe Leu Asn Asn Gly Lys Cys Val Leu Gln Thr His Cys Asp
 3940 3945 3950
 Cys Lys Asp Ala Glu Gly Gly Leu Val Pro Ala Gly Lys Thr Trp Thr
 3955 3960 3965
 Ser Lys Asp Cys Thr Gln Ser Cys Ala Cys Thr Gly Gly Ala Val Gln
 3970 3975 3980
 Cys Gln Asn Phe Gln Cys Pro Leu Gly Thr Tyr Cys Lys Asp Ser Gly
 3985 3990 3995 4000
 Asp Gly Ser Ser Asn Cys Thr Lys Ile His Lys Gly Ala Met Gly Asp
 4005 4010 4015
 Gly Val Leu Met Ala Gly Gly Ile Arg Ala Leu Gln Cys Pro Ala His
 4020 4025 4030
 Ser His Phe Thr Ser Cys Leu Pro Ser Cys Pro Pro Ser Cys Ser Asn
 4035 4040 4045
 Leu Asp Gly Ser Cys Val Glu Ser Asn Phe Lys Ala Pro Ser Val Cys
 4050 4055 4060
 Lys Lys Gly Cys Ile Cys Gln Pro Gly Tyr Leu Leu Asn Asn Asp Lys
 4065 4070 4075 4080
 Cys Val Leu Arg Ile Gln Cys Gly Cys Lys Asp Thr Gln Gly Gly Leu
 4085 4090 4095
 Ile Pro Ala Gly Arg Thr Trp Ile Ser Ser Asp Cys Thr Lys Ser Cys
 4100 4105 4110
 Ser Cys Met Gly Gly Ile Ile Gln Cys Arg Asp Phe Gln Cys Pro Pro
 4115 4120 4125
 Gly Thr Tyr Cys Lys Glu Ser Asn Asp Ser Ser Arg Thr Cys Ala Lys
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 Ile Pro Leu Gln Cys Pro Ala His Ser His Tyr Thr Asn Cys Leu Pro
 4145 4150 4155 4160
 Ala Cys Ser Arg Ser Cys Thr Asp Leu Asp Gly His Cys Glu Gly Thr
 4165 4170 4175
 Ser Pro Lys Val Pro Ser Pro Cys Lys Glu Gly Cys Leu Cys Gln Pro
 4180 4185 4190
 Gly Tyr Val Val His Asn His Lys Cys Val Leu Gln Ile His Cys Gly
 4195 4200 4205

Cys Lys Asp Ala Gln Gly Gly Phe Val Pro Ala Gly Lys Thr Trp Ile
 4210 4215 4220
 Ser Arg Gly Cys Thr Gln Ser Cys Ala Cys Val Gly Gly Ala Val Gln
 4225 4230 4235 4240
 Cys His Asn Phe Thr Cys Pro Thr Gly Thr Gln Cys Gln Asn Ser Ser
 4245 4250 4255
 Cys Ser Lys Ile Thr Val Gln Cys Pro Ala His Ser Gln Tyr Thr Thr
 4260 4265 4270
 Cys Leu Pro Ser Cys Leu Pro Ser Cys Phe Asp Pro Glu Gly Leu Cys
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 Gly Gly Ala Ser Pro Arg Ala Pro Ser Thr Cys Arg Glu Gly Cys Val
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 4305 4310 4315 4320
 Gln Cys Gly Cys Lys Asp Ala Gln Gly Asp Leu Ile Pro Ala Asn Lys
 4325 4330 4335
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 Lys Asp Lys Cys Val Pro Arg Thr Gln Cys Gly Cys Lys Asp Ser Gln
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 Gln Arg Cys Thr Cys Thr Gly Gly Leu Val Gln Cys His Asp Phe Gln
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 Cys Pro Ser Gly Ala Glu Cys Gln Asp Ile Glu Asp Gly Asn Ser Asn
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 Cys Val Glu Ile Thr Val Gln Cys Pro Ala His Ser His Tyr Ser Lys
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Cys Glu Pro Asp Tyr Val Leu Ser Asn Asp Lys Cys Val Pro Ser Ser
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 Glu Cys Gly Cys Lys Asp Ala His Gly Val Leu Ile Pro Glu Ser Lys
 4565 4570 4575
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 4610 4615 4620
 Pro Ala His Ser Leu Tyr Thr Asn Cys Leu Pro Ser Cys Leu Pro Ser
 4625 4630 4635 4640
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 Ser Thr Cys Lys Glu Gly Cys Ile Cys Gln Ser Gly Tyr Val Leu His
 4660 4665 4670
 Lys Asn Lys Cys Met Leu Arg Ile His Cys Asp Cys Lys Asp Phe Gln
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 4785 4790 4795 4800
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 4945 4950 4955 4960
 Met Leu Arg Ala Gln Arg Leu Leu Leu Val Thr Asp Phe Glu Met Val
 4965 4970 4975
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 4980 4985 4990
 Tyr Arg Gly Leu Thr Arg Gly Leu Cys Gly Asn Tyr Asp Arg Asp Gln
 4995 5000 5005
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 5010 5015 5020
 Val Phe Gly Asn Ser Trp Glu Val Lys Ala Gln His Ala Phe Phe Arg
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 Phe Pro Arg Ala Leu Pro Glu Asp Glu Glu Arg Asp Glu Glu Pro Asp
 5045 5050 5055
 Leu Leu Gln Ser Glu Cys Ser Gln Glu Gln Thr Ala Leu Ile Ser Ser
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 Thr Gln Ala Cys Arg Val Leu Val Asp Pro Gln Gly Pro Phe Ala Ala
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 Cys His Gln Ile Ile Ala Pro Glu Pro Phe Glu Gln Arg Cys Met Leu
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 Asp Met Cys Thr Gly Trp Lys Thr Lys Glu Glu Glu Glu Leu Arg Cys
 5105 5110 5115 5120
 Arg Val Leu Ser Gly Tyr Ala Ile Ile Cys Gln Glu Ala Gly Ala Asn
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 5185 5190 5195 5200
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 5205 5210 5215

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Ser Phe Val Thr Asn Asp Cys Ser Gln His Cys Thr Cys Ala Ser Gln
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 <211> 5374
 <212> PRT
 <213> Mus musculus

<400> 75
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 Pro Met Val His Thr Ser Arg Glu Asp Ser Ile Leu Ser Lys Cys Asp
 35 40 45
 Phe Glu Asp Asn Ser Arg Pro Phe Cys Asp Trp Ser Gln Met Ser Ala
 50 55 60
 Asp Asp Gly Asp Trp Ile Arg Thr Thr Gly Pro Ser Leu Thr Gly Thr
 65 70 75 80
 Ser Gly Pro Pro Gly Gly Tyr Pro Asn Gly Glu Gly Tyr Tyr Leu His
 85 90 95
 Met Asp Pro Lys Thr Phe Pro Gln Gly Gly Val Ala Arg Leu Arg Ser
 100 105 110
 Pro Asp Ile Trp Glu Gln Gly Pro Leu Cys Val His Phe Ala Phe His
 Page 174

115					120					125					
Met	Phe	Gly	Leu	Ser	Trp	Gly	Ala	Gln	Leu	Arg	Leu	Leu	Leu	Leu	Arg
	130					135					140				
Gly	Arg	Lys	His	Leu	Arg	Pro	Tyr	Val	Leu	Trp	Lys	His	Val	Asn	Thr
145					150					155					160
Gln	Ser	Pro	Ser	Trp	Met	Pro	Thr	Thr	Val	Thr	Val	Pro	Ala	Asp	His
				165					170					175	
Asp	Ile	Pro	Ser	Trp	Leu	Met	Phe	Glu	Gly	Met	Arg	Gly	Asn	Thr	Ala
			180					185					190		
Tyr	Leu	Asp	Ile	Ser	Leu	Asp	Gly	Leu	Ser	Ile	Gln	Arg	Gly	Thr	Cys
		195					200					205			
Asn	Gln	Val	Cys	Met	Ser	Gln	Met	Cys	Thr	Phe	Asp	Thr	Leu	Asn	Asp
	210					215					220				
Leu	Cys	Gly	Trp	Ser	Trp	Val	Pro	Thr	Ala	Thr	Gly	Ala	Lys	Trp	Thr
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Gln	Lys	Lys	Gly	Pro	Thr	Gly	Lys	Gln	Gly	Val	Gly	Pro	Ala	Glu	Asp
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Phe	Ser	Asn	Pro	Gly	Asn	Gly	Tyr	Tyr	Met	Leu	Leu	Asp	Ser	Thr	Asn
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Ala	Arg	Pro	Gly	Gln	Lys	Ala	Val	Leu	Leu	Ser	Pro	Leu	Ser	His	Ser
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Arg	Gly	Cys	Met	Thr	Leu	Ser	Phe	His	Tyr	Ile	Met	His	Gly	Gln	Gly
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His	Glu	Glu	Gly	Leu	Phe	Val	Tyr	Ala	Thr	Phe	Leu	Gly	Asn	Ile	Arg
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Lys	Tyr	Thr	Leu	Phe	Ser	Gly	His	Pro	Gly	Pro	Asp	Trp	Gln	Ala	Val
				325					330					335	
Ser	Val	Asn	Tyr	Thr	Gly	Gln	Gly	Gln	Ile	Gln	Phe	Met	Val	Val	Gly
			340					345					350		
Met	Phe	Gly	Asn	Ile	Pro	Glu	Pro	Ala	Ile	Ala	Val	Asp	Ala	Ile	Ser
		355					360					365			
Ile	Ala	Pro	Cys	Gly	Glu	Ser	Phe	Pro	Gln	Cys	Asp	Phe	Glu	Asp	Arg
	370					375					380				
Val	His	Pro	Phe	Cys	Asp	Trp	Asn	Gln	Val	Tyr	Gly	Asp	Met	Gly	His
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Trp	Ser	Trp	Gly	Ser	Lys	Ser	Val	Pro	Thr	Leu	Ile	Ala	Gly	Ser	Pro
				405					410					415	
Arg	Glu	Phe	Pro	Tyr	Gly	Gly	Glu	His	Tyr	Ile	Phe	Phe	Asp	Ser	Val
			420					425					430		
Lys	Leu	Ser	Gln	Glu	Gly	Gln	Ser	Ala	Arg	Leu	Val	Ser	Pro	Pro	Phe
		435					440					445			
Cys	Ala	Pro	Gly	Asp	Ile	Cys	Val	Glu	Phe	Ala	Tyr	His	Met	Tyr	Gly

450 455 460
 Leu Gly Lys Gly Thr Thr Leu Lys Leu Leu Leu Gly Ser Pro Ala Gly
 465 470 475 480
 Ser Phe Pro Ile Pro Leu Trp Asn Arg Val Gly Ser Gln Ser Ser Gly
 485 490 495
 Trp Met Asn Ser Ser Val Thr Ile Pro Lys Gly Tyr Gln Gln Pro Met
 500 505 510
 Gln Leu Phe Ile Glu Ala Thr Arg Gly Thr Ser Thr Ala Phe Val Val
 515 520 525
 Ala Leu Asn Phe Ile Leu Ile Ser His Gly Pro Cys Arg Val Leu Leu
 530 535 540
 Gln Thr Glu Ile Pro Ser Ser Pro Leu Leu Pro Pro Thr Gly Pro Ser
 545 550 555 560
 Glu Ser Thr Val Pro Thr Leu Pro Met Glu Gln Pro Thr Ser Pro Thr
 565 570 575
 Lys Ala Thr Thr Val Thr Ile Glu Ile Pro Thr Thr Pro Thr Glu Glu
 580 585 590
 Ala Thr Ile Pro Thr Glu Thr Thr Thr Val Pro Thr Glu Val Ile Asn
 595 600 605
 Val Ser Pro Lys Glu Thr Ser Ile Pro Pro Glu Val Thr Ile Pro Thr
 610 615 620
 Glu Val Ile Thr Val Ser Pro Glu Glu Ile Ile Ser Pro Thr Glu Val
 625 630 635 640
 Thr Pro Val Pro Thr Asp Val Thr Ala Ala Tyr Val Glu Ala Thr Asn
 645 650 655
 Ala Ser Pro Glu Glu Thr Ser Val Pro Pro Glu Val Thr Ile Leu Thr
 660 665 670
 Glu Val Thr Thr Val Ser Pro Glu Glu Thr Thr Val Pro Thr Glu Val
 675 680 685
 Pro Ile Val Leu Ile Glu Ala Thr Ala Phe Pro Thr Gly Glu Thr Thr
 690 695 700
 Leu Tyr Thr Glu Val Pro Thr Val Pro Thr Glu Val Thr Gly Val His
 705 710 715 720
 Thr Glu Val Thr Asn Val Ser Pro Glu Glu Thr Ser Val Pro Thr Glu
 725 730 735
 Glu Thr Ile Ser Thr Glu Val Thr Thr Val Ser Pro Glu Glu Thr Thr
 740 745 750
 Leu Pro Thr Glu Val Pro Thr Val Ser Thr Glu Val Thr Asn Val Ser
 755 760 765
 Pro Glu Glu Thr Ser Val Pro Pro Glu Glu Thr Ile Leu Thr Glu Ile
 770 775 780
 Thr Thr Val Ser Pro Glu Glu Thr Val Phe Pro Thr Glu Gly Thr Thr
 Page 176

785 790 795 800
 Leu Pro Thr Glu Val₈₀₅ Leu Thr Val Pro Ile₈₁₀ Glu Val Thr Thr Phe₈₁₅ Pro
 Thr Gly Glu Thr₈₂₀ Thr Val Pro Thr Glu₈₂₅ Val Pro Thr Val Ser₈₃₀ Thr Glu
 Met Thr Gly₈₃₅ Val His Thr Glu₈₄₀ Thr Thr Val Phe₈₄₅ Pro Glu Glu Thr
 Ser Ile₈₅₀ Pro Thr Glu Val Ala₈₅₅ Thr Val Leu Pro Ala₈₆₀ Ser Ile Pro Pro
 Glu₈₆₅ Glu Thr Thr Thr₈₇₀ Pro Thr Glu Val Thr₈₇₅ Thr Thr Pro Pro Glu₈₈₀
 Thr Thr Ile Pro Ala₈₈₅ Glu Val Thr Thr Val₈₉₀ Pro Pro Val Ser₈₉₅ Ile Pro
 Ser Glu Glu Thr₉₀₀ Thr Thr Pro Thr Glu₉₀₅ Val Thr Thr Thr Pro₉₁₀ Pro Glu
 Glu Thr Thr₉₁₅ Ile Pro Ala Glu Val₉₂₀ Thr Thr Val Pro₉₂₅ Pro Val Ser Ile
 Pro Ser₉₃₀ Glu Glu Thr Thr₉₃₅ Pro Thr Glu Val Thr₉₄₀ Thr Thr Pro Pro
 Glu₉₄₅ Glu Thr Thr Ile₉₅₀ Pro Ala Glu Val Thr₉₅₅ Thr Val Pro Pro Val Ser₉₆₀
 Ile Pro Ser Glu₉₆₅ Glu Thr Thr Ile Pro Thr₉₇₀ Glu Val Thr Thr Val₉₇₅ Pro
 Pro Glu Glu Thr₉₈₀ Thr Ile Pro Ala Glu₉₈₅ Val Thr Thr Val Pro₉₉₀ Pro Val
 Ser Ile₉₉₅ Pro Ser Glu Glu Thr Thr₁₀₀₀ Ile Pro Thr Glu Val₁₀₀₅ Thr Thr Val
 Pro Pro₁₀₁₀ Glu Glu Thr Thr₁₀₁₅ Ile Pro Ala Glu Val₁₀₂₀ Thr Thr Thr Pro Pro
 Glu₁₀₂₅ Glu Thr Thr Ile₁₀₃₀ Pro Thr Glu Val Thr₁₀₃₅ Thr Val Pro Pro Ala Ser₁₀₄₀
 Ile Pro Pro Glu₁₀₄₅ Glu Thr Ala Ser Leu Thr₁₀₅₀ Glu Val Thr Thr Thr Pro₁₀₅₅
 Pro Glu Glu₁₀₆₀ Thr Thr Pro Thr Glu₁₀₆₅ Val Thr Thr Val Pro₁₀₇₀ Pro Glu
 Lys Thr Thr₁₀₇₅ Ile Pro Thr Glu Val₁₀₈₀ Thr Thr Val Pro₁₀₈₅ Pro Ala Ser Ile
 Phe₁₀₉₀ Pro Glu Glu Thr Thr Val₁₀₉₅ Pro Pro Glu Glu Thr₁₁₀₀ Thr Ile Ala Ser
 Glu₁₁₀₅ Glu Thr Thr Val₁₁₁₀ Ser Thr Gln Glu Thr₁₁₁₅ Thr Leu Leu Thr Glu₁₁₂₀ Gln
 Ser Ala Val Thr Gln Thr Ser Ile Ala Cys Arg Pro Pro Cys Pro Ser
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 Pro Pro Leu Met Pro Ile Gly Pro Leu Leu Ser Lys Pro Pro Gly Val
 1140 1145 1150
 Ser Met Phe Ser Leu Ala Pro Thr Thr Gly Val Ser Thr Thr Glu Ser
 1155 1160 1165
 Cys Pro Pro Asn Ala His Ile Glu Leu Cys Ala Cys Pro Ala Ser Cys
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 Glu Ser Pro Lys Pro Ser Cys Gln Pro Pro Cys Ile Pro Gly Cys Val
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 Cys Asn Pro Gly Phe Leu Phe Ser Asn Asn Gln Cys Ile Asn Glu Ser
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 Arg Leu Ser Trp Gln Val Glu Glu Asp Glu Asp Lys Asp Trp Val Ser
 1445 1450 1455
 Ser Arg Cys Gln Lys Lys Lys Asn Pro Pro Ser Cys Asp Ala Ala Leu
 Page 178

1460 1465 1470
 Gly Ser Thr Met Ser Gly Pro Lys Leu Cys Gly Gln Leu Val Asn Pro
 1475 1480 1485
 Ser Gly Pro Phe Glu Ala Cys Leu Leu His Leu Lys Ala Ser Ser Phe
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 Lys Leu Cys Ala His Met Ser Ala Met Thr Ala Thr Cys Gln Asp Ala
 1525 1530 1535
 Gly Tyr Pro Val Lys Pro Trp Arg Glu Pro Gln Phe Cys Pro Leu Val
 1540 1545 1550
 Cys Pro Lys Asn Ser Arg Tyr Ser Leu Cys Ala Lys Pro Cys Pro Glu
 1555 1560 1565
 Thr Cys His Pro Ile Ser Thr Thr Gln His Cys Ser Asp Lys Cys Val
 1570 1575 1580
 Glu Gly Cys Glu Cys Asp Pro Gly Phe Ile Leu Ser Gly Ser Glu Cys
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 Val Pro Ser Ser Gln Cys Gly Cys Thr Ser Phe Gln Gly Arg Tyr Phe
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 Lys Val Gln Glu Gln Trp Phe Asn Pro Asp Cys Lys Glu Ile Cys Thr
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 Gly Ala Ala Thr Cys Met Val Ser Gly Asp Pro His Tyr Leu Thr Phe
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 Asp Gly Ala Leu His His Phe Met Gly Thr Cys Thr Tyr Val Leu Thr
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 1765 1770 1775
 Thr Asn Phe Gly Leu Gln Val Arg Tyr Asp Gly Arg His Leu Val Glu
 1780 1785 1790
 Val Thr Val Pro Ser Ser Tyr Thr Gly Ser Leu Cys Gly Leu Cys Gly
 Page 179

1795 1800 1805
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 Pro Ala Gly Asn Ser Leu Leu Leu Gly Ala Ala Trp Lys Ile Leu Glu
 1825 1830 1835 1840
 Ala Ser Asp Pro Gly Cys Phe Leu Val Gly Gly Lys Pro Ser Arg Cys
 1845 1850 1855
 Ala Asp Ser Asp Met Asp Asp Val Trp Thr Lys Lys Cys Ala Ile Leu
 1860 1865 1870
 Met Asn Pro Leu Gly Pro Phe Ser Asn Cys His Glu Ala Val Pro Pro
 1875 1880 1885
 Gln Ala Ser Phe Ser Ser Cys Val Tyr Gly Gln Cys Glu Thr Asn Gly
 1890 1895 1900
 Asp Asn Leu Thr Leu Cys His Ser Leu Gln Ala Tyr Ala Ser Leu Cys
 1905 1910 1915 1920
 Ala Gln Ala Gly Gln Val Thr Thr Trp Arg Asn Ser Thr Phe Cys Pro
 1925 1930 1935
 Met Arg Cys Pro Pro Arg Ser Ser Tyr Asn Pro Cys Ala Asn Ser Cys
 1940 1945 1950
 Pro Ala Thr Cys Leu Thr Leu Ser Thr Pro Arg Asp Cys Pro Thr Leu
 1955 1960 1965
 Pro Cys Val Glu Gly Cys Glu Cys Gln Ser Gly His Ile Leu Ser Gly
 1970 1975 1980
 Thr Thr Cys Val Pro Leu Arg Gln Cys Gly Cys Ser Asp Gln Asp Gly
 1985 1990 1995 2000
 Ser Tyr His Leu Leu Gly Glu Ser Trp Tyr Thr Glu Lys Thr Cys Thr
 2005 2010 2015
 Thr Leu Cys Thr Cys Ser Ala His Ser Asn Ile Thr Cys Ser Pro Thr
 2020 2025 2030
 Ala Cys Lys Ala Asn His Val Cys Leu Arg Gln Glu Gly Leu Leu Arg
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 Cys Ala Ala Glu Met Gly Glu Cys Arg Ile Ser Glu Asp Ser Gln Ile
 2050 2055 2060
 Val Ser Phe Asp Asp His Ser His Pro Ile Gln Asp Thr Cys Thr Tyr
 2065 2070 2075 2080
 Ile Leu Val Lys Val Cys His Pro Asn Thr Asn Met Pro Phe Phe Met
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 Gly Val Tyr Gln Leu Tyr Ile Asp Ile Phe Asn Phe His Ile Thr Leu
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 Gln Lys Asp His Leu Val Leu Ile Ser Leu Ile Asn Asp Ser Ile Val
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Lys Val Cys Gly Val Cys Gly Asn Phe Asn Gly Glu Glu Glu Asp Glu
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Leu Met Thr Pro Ser Gly Glu Leu Ala Glu Asp Glu Gln Glu Phe Met
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2275 2280 2285
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Gln Gly Leu Lys Pro Pro Ile Trp Arg Asn Ser Ser Phe Cys Pro Leu
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Glu Cys Pro Ala His Ser His Tyr Thr Asn Cys Leu Pro Ser Cys Pro
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Pro Ser Cys Leu Asp Pro Asp Ser Arg Cys Glu Gly Ser Gly His Lys
2355 2360 2365
Val Pro Ala Thr Cys Arg Glu Gly Cys Ile Cys Gln Pro Asp Tyr Val
2370 2375 2380
Leu Leu Asn Asp Lys Cys Val Leu Arg Ser His Cys Gly Cys Lys Asp
2385 2390 2395 2400
Ala Gln Gly Val Phe Ile Pro Ala Gly Lys Thr Trp Ile Ser Glu Asp
2405 2410 2415
Cys Thr Gln Ser Cys Thr Cys Met Lys Gly Ser Met Arg Cys Trp Asp
2420 2425 2430
Phe Gln Cys Pro Pro Gly Thr Tyr Cys Lys Asn Ser Asn Asp Gly Ser
2435 2440 2445
Ser Asn Cys Val Lys Ile Ser Leu Gln Cys Pro Ala His Ser Lys Phe
2450 2455 2460
Thr Asp Cys Leu Pro Pro Cys His Pro Ser Cys Ser Asp Pro Asp Gly
Page 181

2465 2470 2475 2480
 His Cys Glu Gly Ile Ser Thr Asn Ala His Ser Asn Cys Lys Glu Gly
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 Cys Val Cys Gln Pro Gly Tyr Val Leu Arg Asn Asp Lys Cys Val Leu
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 Gly Gly Ala Ile Gln Cys Gln Asn Phe Lys Cys Pro Ser Glu Ala Tyr
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 Cys Gln Asp Leu Glu Asp Gly Asn Ser Asn Cys Thr Ser Ile Pro Leu
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2805 2810 2815
 Gln Cys Pro Ala His Ser His Tyr Thr Asn Cys Leu Pro Thr Cys Gln
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 Pro Ser Cys Ser Asp Pro Asp Gly His Cys Glu Gly Ser Ser Thr Lys
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 2865 2870 2875 2880
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 2885 2890 2895
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 Tyr His Cys Ser Ser Gly Thr Tyr Cys Lys Asp Met Glu Asp Asp Ser
 2915 2920 2925
 Ser Ser Cys Ala Thr Ile Thr Leu Gln Cys Pro Ala His Ser His Phe
 2930 2935 2940
 Thr Asn Cys Leu Pro Pro Cys Gln Pro Ser Cys Leu Asp Ser Glu Gly
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 His Cys Glu Gly Ser Thr Thr Lys Ala Pro Ser Ala Cys Gln Glu Gly
 2965 2970 2975
 Cys Val Cys Glu Pro Asp Tyr Val Val Leu Asn Asn Lys Cys Val Pro
 2980 2985 2990
 Arg Ile Glu Cys Gly Cys Lys Asp Ala Gln Gly Val Leu Ile Pro Ala
 2995 3000 3005
 Asp Lys Thr Trp Ile Asn Arg Gly Cys Thr Gln Ser Cys Thr Cys Lys
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 Gly Gly Ala Ile Gln Cys Gln Lys Phe Gln Cys Pro Ser Glu Thr Tyr
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 Cys Lys Asp Ile Glu Asp Gly Asn Ser Asn Cys Thr Arg Ile Ser Leu
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 Gln Cys Pro Ala Asn Ser Asn Phe Thr Ser Cys Leu Pro Ser Cys Gln
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 Pro Ser Cys Ser Asn Thr Asp Val His Cys Glu Gly Ser Ser Pro Asn
 3075 3080 3085
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 3090 3095 3100
 Leu His Asn Asp Lys Cys Ile Leu Arg Asn Gln Cys Gly Cys Lys Asp
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 Ala Gln Gly Ala Leu Ile Pro Glu Gly Lys Thr Trp Ile Thr Ser Gly
 3125 3130 3135
 Cys Thr Gln Ser Cys Asn Cys Thr Gly Gly Ala Ile Gln Cys Gln Asn
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3140 3145 3150
 Phe Gln Cys Pro Leu Lys Thr Tyr Cys Lys Asp Leu Lys Asp Gly Ser
 3155 3160 3165
 Ser Asn Cys Thr Asn Ile Pro Leu Gln Cys Pro Ala His Ser Arg Tyr
 3170 3175 3180
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 Cys Ile Cys Gln Pro Gly Tyr Leu Met His Lys Asn Lys Cys Val Leu
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 Asp Lys Thr Trp Ile Ser Arg Gly Cys Thr Gln Ser Cys Thr Cys Ser
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 Ala Gly Ala Ile His Cys Arg Asn Phe Lys Cys Pro Ser Gly Thr Tyr
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 Ser Cys Ala Cys Val Glu Gly Asn Ile Gln Cys Gln Asn Phe Gln Cys
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 Pro Pro Glu Thr Tyr Cys Lys Asp Asn Ser Glu Gly Ser Ser Thr Cys
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 Page 184

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 Thr Cys Lys Glu Gly Cys Ile Cys Gln Ser Gly Tyr Val Leu Ser Asn
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 Ala Leu Ile Pro Glu Asp Lys Thr Trp Val Ser Arg Gly Cys Thr Gln
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 Ser Cys Val Cys Thr Gly Gly Ser Ile Gln Cys Leu Ser Phe Gln Cys
 3620 3625 3630
 Pro Pro Gly Ala Tyr Cys Lys Asp Asn Glu Asp Gly Ser Ser Asn Cys
 3635 3640 3645
 Ala Arg Ile Pro Pro Gln Cys Pro Ala Asn Ser His Tyr Thr Asp Cys
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 Phe Pro Pro Cys Pro Pro Ser Cys Ser Asp Pro Glu Gly His Cys Glu
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 Ala Ser Gly Pro Arg Val Pro Ser Thr Cys Arg Glu Gly Cys Leu Cys
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 Asn Pro Gly Phe Val Leu Asp Arg Asp Lys Cys Val Pro Arg Val Glu
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 Page 185

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 Pro Pro Gly Thr Tyr Cys Asn His Asn Asn Asn Cys Ala Lys Ile Pro
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 Page 186

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 Lys Val Pro Ser Pro Cys Lys Glu Gly Cys Leu Cys Gln Pro Gly Tyr
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Gln Cys His Asp Phe Ser Cys Pro Thr Gly Ser Arg Cys Leu Asp Asn
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Asn Glu Gly Asn Ser Asn Cys Val Thr Tyr Ala Leu Lys Cys Pro Ala
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His Ser Leu Tyr Thr Asn Cys Leu Pro Ser Cys Leu Pro Ser Cys Ser
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Asp Pro Glu Gly Leu Cys Gly Gly Thr Ser Pro Glu Val Pro Ser Thr
4645 4650 4655

Cys Lys Glu Gly Cys Ile Cys Gln Ser Gly Tyr Val Leu His Lys Asn
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Lys Cys Met Leu Arg Ile His Cys Asp Cys Lys Asp Phe Gln Gly Ser
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Leu Ile Lys Thr Gly Gln Thr Trp Ile Ser Ser Gly Cys Ser Lys Ile
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Thr Ser His Lys Ala Pro Ser Thr Cys Arg Glu Gly Cys Val Cys Gln
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Pro Gly Tyr Leu Leu Asn Lys Asp Thr Cys Val His Lys Asn Gln Cys
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 Ile Phe Gly Asp Pro Tyr Tyr Leu Thr Phe Asp Gly Phe Thr Tyr His
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 Page 189

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 5365 5370

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 <211> 2601
 <212> PRT
 <213> Homo sapiens

<400> 76
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 Phe Arg Lys Glu Lys Pro Pro Asp Gln Lys Leu Val Val Arg Ser Ser
 20 25 30
 Arg Asp Asn Tyr Val Leu Thr Gln Cys Asp Phe Glu Asp Asp Ala Lys
 35 40 45
 Pro Leu Cys Asp Trp Ser Gln Val Ser Ala Asp Asp Glu Asp Trp Val
 50 55 60
 Arg Ala Ser Gly Pro Ser Pro Thr Gly Ser Thr Gly Ala Pro Gly Gly
 65 70 75 80

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Tyr	Pro	Asn	Gly	Glu ₈₅	Gly	Ser	Tyr	Leu	His ₉₀	Met	Glu	Ser	Asn	Ser ₉₅	Phe
His	Arg	Gly	Gly ₁₀₀	Val	Ala	Arg	Leu	Leu ₁₀₅	Ser	Pro	Asp	Leu	Trp ₁₁₀	Glu	Gln
Gly	Pro	Leu ₁₁₅	Cys	Val	His	Phe	Ala ₁₂₀	His	His	Met	Phe	Gly ₁₂₅	Leu	Ser	Trp
Gly	Ala	Gln	Leu	Arg	Leu	Leu ₁₃₅	Leu	Leu	Ser	Gly	Glu ₁₄₀	Glu	Gly	Arg	Arg
Pro	Asp	Val	Leu	Trp	Lys ₁₅₀	His	Trp	Asn	Thr	Gln ₁₅₅	Arg	Pro	Ser	Trp	Met ₁₆₀
Leu	Thr	Thr	Val	Thr ₁₆₅	Val	Pro	Ala	Gly	Phe ₁₇₀	Thr	Leu	Pro	Thr	Arg ₁₇₅	Leu
Met	Phe	Glu	Gly ₁₈₀	Thr	Arg	Gly	Ser	Thr ₁₈₅	Ala	Tyr	Leu	Asp	Ile ₁₉₀	Ala	Leu
Asp	Ala	Leu ₁₉₅	Ser	Ile	Arg	Arg	Gly ₂₀₀	Ser	Cys	Asn	Arg	Val ₂₀₅	Cys	Met	Met
Gln	Thr ₂₁₀	Cys	Ser	Phe	Asp	Ile ₂₁₅	Pro	Asn	Asp	Leu	Cys ₂₂₀	Asp	Trp	Thr	Trp
Ile	Pro	Thr	Ala	Ser	Gly ₂₃₀	Ala	Lys	Trp	Thr	Gln ₂₃₅	Lys	Lys	Gly	Ser	Ser ₂₄₀
Gly	Lys	Pro	Gly	Val ₂₄₅	Gly	Pro	Asp	Gly	Asp ₂₅₀	Phe	Ser	Ser	Pro	Gly ₂₅₅	Ser
Gly	Cys	Tyr	Met ₂₆₀	Leu	Leu	Asp	Pro	Lys ₂₆₅	Asn	Ala	Arg	Pro	Gly ₂₇₀	Gln	Lys
Ala	Val	Leu ₂₇₅	Leu	Ser	Pro	Val	Ser ₂₈₀	Leu	Ser	Ser	Gly	Cys ₂₈₅	Leu	Ser	Phe
Ser	Phe	His	Tyr	Ile	Leu	Arg ₂₉₅	Gly	Gln	Ser	Pro	Gly ₃₀₀	Ala	Ala	Leu	His
Ile	Tyr	Ala	Ser	Val	Leu ₃₁₀	Gly	Ser	Ile	Arg	Lys ₃₁₅	His	Thr	Leu	Phe	Ser ₃₂₀
Gly	Gln	Pro	Gly	Pro ₃₂₅	Asn	Trp	Gln	Ala	Val ₃₃₀	Ser	Val	Asn	Tyr	Thr ₃₃₅	Ala
Val	Gly	Arg	Ile ₃₄₀	Gln	Phe	Ala	Val	Val ₃₄₅	Gly	Val	Phe	Gly	Lys ₃₅₀	Thr	Pro
Glu	Pro	Ala	Val ₃₅₅	Ala	Val	Asp	Ala ₃₆₀	Thr	Ser	Ile	Ala	Pro	Cys	Gly	Glu
Gly	Phe	Pro	Gln	Cys	Asp	Phe ₃₇₅	Glu	Asp	Asn	Ala	His ₃₈₀	Pro	Phe	Cys	Asp
Trp	Val	Gln	Thr	Ser	Gly ₃₉₀	Asp	Gly	Gly	His	Trp ₃₉₅	Ala	Leu	Gly	His	Lys ₄₀₀
Asn	Gly	Pro	Val	His ₄₀₅	Gly	Met	Gly	Pro	Ala ₄₁₀	Gly	Gly	Phe	Pro	Asn	Ala ₄₁₅

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Gly Gly His Tyr Ile Tyr Leu Glu Ala Asp Glu Phe Ser Gln Ala Gly
420 425 430

Gln Ser Val Arg Leu Val Ser Arg Pro Phe Cys Ala Pro Gly Asp Ile
435 440 445

Cys Val Glu Phe Ala Tyr His Met Tyr Gly Leu Gly Glu Gly Thr Met
450 455 460

Leu Glu Leu Leu Leu Gly Ser Pro Ala Gly Ser Pro Pro Ile Pro Leu
465 470 475 480

Trp Lys Arg Val Gly Ser Gln Arg Pro Tyr Trp Gln Asn Thr Ser Val
485 490 495

Thr Val Pro Ser Gly His Gln Gln Pro Met Gln Leu Ile Phe Lys Gly
500 505 510

Ile Gln Gly Ser Asn Thr Ala Ser Val Val Ala Met Gly Phe Ile Leu
515 520 525

Ile Asn Pro Gly Thr Cys Pro Val Lys Val Leu Pro Glu Leu Pro Pro
530 535 540

Val Ser Pro Val Ser Ser Thr Gly Pro Ser Glu Thr Thr Gly Leu Thr
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Glu Asn Pro Thr Ile Ser Thr Lys Lys Pro Thr Val Ser Ile Glu Lys
565 570 575

Pro Ser Val Thr Thr Glu Lys Pro Thr Val Pro Lys Glu Lys Pro Thr
580 585 590

Ile Pro Thr Glu Lys Pro Thr Ile Ser Thr Glu Lys Pro Thr Ile Pro
595 600 605

Ser Glu Lys Pro Asn Met Pro Ser Glu Lys Pro Thr Ile Pro Ser Glu
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Lys Pro Thr Ile Leu Thr Glu Lys Pro Thr Ile Pro Ser Glu Lys Pro
625 630 635 640

Thr Ile Pro Ser Glu Lys Pro Thr Ile Ser Thr Glu Lys Pro Thr Val
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Pro Thr Glu Glu Pro Thr Thr Pro Thr Glu Glu Thr Thr Thr Ser Met
660 665 670

Glu Glu Pro Val Ile Pro Thr Glu Lys Pro Ser Ile Pro Thr Glu Lys
675 680 685

Pro Ser Ile Pro Thr Glu Lys Pro Thr Ile Ser Met Glu Glu Thr Ile
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Ile Ser Thr Glu Lys Pro Thr Ile Ser Pro Glu Lys Pro Thr Ile Pro
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Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Ser Thr Ile Ser Pro Glu
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Lys Pro Thr Thr Pro Thr Glu Lys Pro Thr Ile Pro Thr Glu Lys Pro
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 Pro Thr Glu Lys Leu Thr Ala Leu Arg Pro Pro His Pro Ser Pro Thr
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 Ala Val His Val Thr Val Phe Asp Leu Ser Ile Ser Leu Leu Arg Gly
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 Cys Lys Val Met Leu Asn Gly His Arg Val Ala Leu Pro Val Trp Leu
 1620 1625 1630
 Ala Gln Gly Arg Val Thr Ile Arg Leu Ser Ser Asn Leu Val Leu Leu
 1635 1640 1645
 Tyr Thr Asn Phe Gly Leu Gln Val Arg Tyr Asp Gly Ser His Leu Val
 1650 1655 1660
 Glu Val Thr Val Pro Ser Ser Tyr Gly Gly Gln Leu Cys Gly Leu Cys
 1665 1670 1675 1680
 Gly Asn Tyr Asn Asn Asn Ser Leu Asp Asp Asn Leu Arg Pro Asp Arg
 1685 1690 1695
 Lys Leu Ala Gly Asp Ser Met Gln Leu Gly Ala Ala Trp Lys Leu Pro
 1700 1705 1710
 Glu Ser Ser Glu Pro Gly Cys Phe Leu Val Gly Gly Lys Pro Ser Ser
 1715 1720 1725
 Cys Gln Glu Asn Ser Met Ala Asp Ala Trp Asn Lys Asn Cys Ala Ile
 1730 1735 1740
 Leu Ile Asn Pro Gln Gly Pro Phe Ser Gln Cys His Gln Val Val Pro
 1745 1750 1755 1760

Pro Gln Ser Ser Phe Ala Ser Cys Val His Gly Gln Cys Gly Thr Lys
 1765 1770 1775
 Gly Asp Thr Thr Ala Leu Cys Arg Ser Leu Gln Ala Tyr Ala Ser Leu
 1780 1785 1790
 Cys Ala Gln Ala Gly Gln Ala Pro Ala Trp Arg Asn Arg Thr Phe Cys
 1795 1800 1805
 Pro Met Arg Cys Pro Pro Gly Ser Ser Tyr Ser Pro Cys Ser Ser Pro
 1810 1815 1820
 Cys Pro Asp Thr Cys Ser Ser Ile Asn Asn Pro Arg Asp Cys Pro Lys
 1825 1830 1835 1840
 Ala Leu Pro Cys Ala Glu Ser Cys Glu Cys Gln Lys Gly His Ile Leu
 1845 1850 1855
 Ser Gly Thr Ser Cys Val Pro Leu Gly Gln Cys Gly Cys Thr Asp Pro
 1860 1865 1870
 Ala Gly Ser Tyr His Pro Val Gly Glu Arg Trp Tyr Thr Glu Asn Thr
 1875 1880 1885
 Cys Thr Arg Leu Cys Thr Cys Ser Val His Asn Asn Ile Thr Cys Phe
 1890 1895 1900
 Gln Ser Thr Cys Lys Pro Asn Gln Ile Cys Trp Ala Leu Asp Gly Leu
 1905 1910 1915 1920
 Leu Arg Cys Arg Ala Ser Gly Val Gly Val Cys Gln Leu Pro Gly Glu
 1925 1930 1935
 Ser His Tyr Val Ser Phe Asp Gly Ser Asn His Ser Ile Pro Asp Ala
 1940 1945 1950
 Cys Thr Leu Val Leu Val Lys Val Cys His Pro Ala Met Ala Leu Pro
 1955 1960 1965
 Phe Phe Lys Ile Ser Ala Lys His Glu Lys Glu Glu Gly Gly Thr Glu
 1970 1975 1980
 Ala Phe Arg Leu His Glu Val Tyr Ile Asp Ile Tyr Asp Ala Gln Val
 1985 1990 1995 2000
 Thr Leu Gln Lys Gly His Arg Val Leu Ile Asn Ser Lys Gln Val Thr
 2005 2010 2015
 Leu Pro Ala Ile Ser Gln Ile Pro Gly Val Ser Val Lys Ser Ser Ser
 2020 2025 2030
 Ile Tyr Ser Ile Val Asn Ile Lys Ile Gly Val Gln Val Lys Phe Asp
 2035 2040 2045
 Gly Asn His Leu Leu Glu Ile Glu Ile Pro Thr Thr Tyr Tyr Gly Lys
 2050 2055 2060
 Val Cys Gly Met Cys Gly Asn Phe Asn Asp Glu Glu Glu Asp Glu Leu
 2065 2070 2075 2080
 Met Met Pro Ser Asp Glu Val Ala Asn Ser Asp Ser Glu Phe Val Asn
 2085 2090 2095

Ser Trp Lys Asp Lys Asp Ile Asp Pro Ser Cys Gln Ser Leu Leu Val
 2100 2105 2110
 Asp Glu Gln Gln Ile Pro Ala Glu Gln Gln Glu Asn Pro Ser Gly Asn
 2115 2120 2125
 Cys Arg Ala Ala Asp Leu Arg Arg Ala Arg Glu Lys Cys Glu Ala Ala
 2130 2135 2140
 Leu Arg Ala Pro Val Trp Ala Gln Cys Ala Ser Arg Ile Asp Leu Thr
 2145 2150 2155 2160
 Pro Phe Leu Val Asp Cys Ala Asn Thr Leu Cys Glu Phe Gly Gly Leu
 2165 2170 2175
 Tyr Gln Ala Leu Cys Gln Ala Leu Gln Ala Phe Gly Ala Thr Cys Gln
 2180 2185 2190
 Ser Gln Gly Leu Lys Pro Pro Leu Trp Arg Asn Ser Ser Phe Cys Pro
 2195 2200 2205
 Leu Glu Cys Pro Ala Tyr Ser Ser Tyr Thr Asn Cys Leu Pro Ser Cys
 2210 2215 2220
 Ser Pro Ser Cys Trp Asp Leu Asp Gly Arg Cys Glu Gly Ala Lys Val
 2225 2230 2235 2240
 Pro Ser Ala Cys Ala Glu Gly Cys Ile Cys Gln Pro Gly Tyr Val Leu
 2245 2250 2255
 Ser Glu Asp Lys Cys Val Pro Arg Ser Gln Cys Gly Cys Lys Asp Ala
 2260 2265 2270
 His Gly Gly Ser Ile Pro Leu Gly Lys Ser Trp Val Ser Ser Gly Cys
 2275 2280 2285
 Thr Glu Lys Cys Val Cys Thr Gly Gly Ala Ile Gln Cys Gly Asp Phe
 2290 2295 2300
 Arg Cys Pro Ser Gly Ser His Cys Gln Leu Thr Ser Asp Asn Ser Asn
 2305 2310 2315 2320
 Ser Asn Cys Val Ser Asp Lys Ser Glu Gln Cys Ser Val Tyr Gly Asp
 2325 2330 2335
 Pro Arg Tyr Leu Thr Phe Asp Gly Phe Ser Tyr Arg Leu Gln Gly Arg
 2340 2345 2350
 Met Thr Tyr Val Leu Ile Lys Thr Val Asp Val Leu Pro Glu Gly Val
 2355 2360 2365
 Glu Pro Leu Leu Val Glu Gly Arg Asn Lys Met Asp Pro Pro Arg Ser
 2370 2375 2380
 Ser Ile Phe Leu Gln Glu Val Ile Thr Thr Val Tyr Gly Tyr Lys Val
 2385 2390 2395 2400
 Gln Leu Gln Ala Gly Leu Glu Leu Val Val Asn Asn Gln Lys Met Ala
 2405 2410 2415
 Val Pro Tyr Arg Pro Asn Glu His Leu Arg Val Thr Leu Trp Gly Gln
 2420 2425 2430

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Arg Leu Tyr Leu Val Thr Asp Phe Glu Leu Val Val Ser Phe Gly Gly
 2435 2440 2445

Arg Lys Asn Ala Val Ile Ser Leu Pro Ser Met Tyr Glu Gly Leu Val
 2450 2455 2460

Ser Gly Leu Cys Gly Asn Tyr Asp Lys Asn Arg Lys Asn Asp Met Met
 2465 2470 2475 2480

Leu Pro Ser Gly Ala Leu Thr Gln Asn Leu Asn Thr Phe Gly Asn Ser
 2485 2490 2495

Trp Glu Val Lys Thr Glu Asp Ala Leu Leu Arg Phe Pro Arg Ala Ile
 2500 2505 2510

Pro Ala Glu Glu Glu Gly Gln Gly Ala Glu Leu Gly Leu Arg Thr Gly
 2515 2520 2525

Leu Gln Val Ser Glu Cys Ser Pro Glu Gln Leu Ala Ser Asn Ser Thr
 2530 2535 2540

Gln Ala Cys Arg Val Leu Ala Asp Pro Gln Gly Pro Phe Ala Ala Cys
 2545 2550 2555 2560

His Gln Thr Val Ala Pro Glu Pro Phe Gln Glu His Cys Val Leu Asp
 2565 2570 2575

Leu Cys Ser Ala Gln Asp Pro Arg Glu Gln Glu Glu Leu Arg Cys Gln
 2580 2585 2590

Val Leu Ser Gly Trp Ala Ala Ala Phe
 2595 2600

<210> 77

<211> 170

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: MAM domain
 sequence

<400> 77

Cys Asp Phe Glu Asp Gly Ser His Pro Phe Cys Gly Trp Ser Gln Asp
 1 5 10 15

Ser Gly Asp Asp Gly Asp Asp Leu Gln Trp Thr Arg Val Asn Ser Ala
 20 25 30

Thr Gly Gly Ser Thr Gly Pro Arg Gly Asp His Thr Thr Gly Asn Gly
 35 40 45

His Tyr Met Tyr Val Asp Thr Ser Ser Gly Leu Leu Gln Glu Gly Gln
 50 55 60

Lys Ala Arg Leu Leu Ser Pro Pro Leu Pro Pro Asn Arg Ser Pro Glu
 65 70 75 80

Cys Cys Leu Thr Phe Trp Tyr His Met Tyr Gly Ser Gly Val Gly Thr
 85 90 95

Pro Gly Leu Asn Val Tyr Val Arg Glu Asn Gly Glu Thr Leu Leu Trp
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CORAZZ21:APP															
100							105					110			
Ser	Arg	Ser	Gly	His	Gln	Gly	Gly	Gln	Trp	Leu	Leu	Ala	Glu	Val	Thr
		115					120					125			
Leu	Pro	Thr	Phe	Ser	Thr	Lys	Pro	Phe	Gln	Val	Val	Phe	Glu	Gly	Thr
	130					135					140				
Arg	Gly	Gly	Gly	Ser	Arg	Gly	Gly	Ile	Ala	Leu	Asp	Asp	Ile	Ser	Leu
145					150					155					160
Ser	Thr	His	Ile	Glu	Gly	Pro	Cys	Asn	Gln						
				165					170						

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<210> 78
<211> 170
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: MAM domain
sequence

<400>	78															
Cys 1	Asp	Phe	Glu	Asp 5	Gly	Ser	His	Pro	Phe 10	Cys	Gly	Trp	Ser	Gln 15	Asp	
Ser	Gly	Asp	Asp 20	Gly	Asp	Asp	Leu	Gln 25	Trp	Thr	Arg	Val	Asn 30	Ser	Ala	
Thr	Gly	Gly 35	Ser	Thr	Gly	Pro	Arg 40	Gly	Asp	His	Thr	Thr 45	Gly	Asn	Gly	
His	Tyr 50	Met	Tyr	Val	Asp	Thr 55	Ser	Ser	Gly	Leu	Leu 60	Gln	Glu	Gly	Gln	
Lys 65	Ala	Arg	Leu	Leu	Ser 70	Pro	Pro	Leu	Pro	Pro 75	Asn	Arg	Ser	Pro	Glu 80	
Cys	Cys	Leu	Thr	Phe 85	Trp	Tyr	His	Met	Tyr 90	Gly	Ser	Gly	Val	Gly 95	Thr	
Pro	Gly	Leu	Asn 100	Val	Tyr	Val	Arg	Glu 105	Asn	Gly	Glu	Thr	Leu 110	Leu	Trp	
Ser	Arg	Ser 115	Gly	His	Gln	Gly	Gly 120	Gln	Trp	Leu	Leu	Ala 125	Glu	Val	Thr	
Leu	Pro 130	Thr	Phe	Ser	Thr	Lys 135	Pro	Phe	Gln	Val	Val 140	Phe	Glu	Gly	Thr	
Arg 145	Gly	Gly	Gly	Ser	Arg 150	Gly	Gly	Ile	Ala	Leu 155	Asp	Asp	Ile	Ser	Leu 160	
Ser	Thr	His	Ile 165	Glu	Gly	Pro	Cys	Asn	Gln 170							

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<210> 79
<211> 812
<212> PRT
<213> Homo sapiens
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<400> 79

Met Gly Trp Arg Pro Arg Arg Ala Arg Gly Thr Pro Leu Leu Leu Leu
 1 5 10 15
 Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln
 20 25 30
 Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
 35 40 45
 Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
 50 55 60
 Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
 65 70 75 80
 Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
 85 90 95
 Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Asp
 100 105 110
 His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val
 115 120 125
 Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg
 130 135 140
 Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp
 145 150 155 160
 Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys
 165 170 175
 Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser
 180 185 190
 Leu Pro Gly Gly Pro Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr
 195 200 205
 Arg Lys Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu
 210 215 220
 Thr Arg His Arg Asn Leu Asn His Thr Lys Gln Arg Leu Leu Glu Val
 225 230 235 240
 Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala
 245 250 255
 Leu Thr Gly Leu Glu Val Trp Thr Glu Arg Asp Arg Ser Arg Val Thr
 260 265 270
 Gln Asp Ala Asn Ala Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg Gly
 275 280 285
 Leu Trp Ala Gln Arg Pro His Asp Ser Ala Gln Leu Leu Thr Gly Arg
 290 295 300
 Ala Phe Gln Gly Ala Thr Val Gly Leu Ala Pro Val Glu Gly Met Cys
 305 310 315 320
 Arg Ala Glu Ser Ser Gly Gly Val Ser Thr Asp His Ser Glu Leu Pro

325 335
Ile Gly Ala Ala Ala Thr Met Ala His Glu Ile Gly His Ser Leu Gly
340 345 350
Leu Ser His Asp Pro Asp Gly Cys Cys Val Glu Ala Ala Ala Glu Ser
355 360 365
Gly Gly Cys Val Met Ala Ala Ala Thr Gly His Pro Phe Pro Arg Val
370 380
Phe Ser Ala Cys Ser Arg Arg Gln Leu Arg Ala Phe Phe Arg Lys Gly
385 390 395 400
Gly Gly Ala Cys Leu Ser Asn Ala Pro Asp Pro Gly Leu Pro Val Pro
405 410 415
Pro Ala Leu Cys Gly Asn Gly Phe Val Glu Ala Gly Glu Glu Cys Asp
420 425 430
Cys Gly Pro Gly Gln Glu Cys Arg Asp Leu Cys Cys Phe Ala His Asn
435 440 445
Cys Ser Leu Arg Pro Gly Ala Gln Cys Ala His Gly Asp Cys Cys Val
450 455 460
Arg Cys Leu Leu Lys Pro Ala Gly Ala Leu Cys Arg Gln Ala Met Gly
465 470 475 480
Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly Thr Ser Ser His Cys Pro
485 490 495
Pro Asp Val Tyr Leu Leu Asp Gly Ser Pro Cys Ala Arg Gly Ser Gly
500 505 510
Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys Gln Gln
515 520 525
Leu Trp Gly Pro Gly Ser His Pro Ala Pro Glu Ala Cys Phe Gln Val
530 535 540
Val Asn Ser Ala Gly Asp Ala His Gly Asn Cys Gly Gln Asp Ser Glu
545 550 555 560
Gly His Phe Leu Pro Cys Ala Gly Arg Asp Ala Leu Cys Gly Lys Leu
565 570 575
Gln Cys Gln Gly Gly Lys Pro Ser Leu Leu Ala Pro His Met Val Pro
580 585 590
Val Asp Ser Thr Val His Leu Asp Gly Gln Glu Val Thr Cys Arg Gly
595 600 605
Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu
610 615 620
Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg
625 630 635 640
Arg Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala
645 650 655
Cys His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala

660 665 670
 Pro Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser
 675 680 685
 Met Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu
 690 695 700
 Ala Met Leu Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu
 705 710 715 720
 Ala Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser
 725 730 735
 Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro
 740 745 750
 His Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro
 755 760 765
 Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu
 770 775 780
 Pro Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro
 785 790 795 800
 Gln Asp Gln Val Gln Met Pro Arg Ser Cys Leu Trp
 805 810

<210> 80
 <211> 728
 <212> PRT
 <213> Homo sapiens

<400> 80
 Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His Tyr Gly Pro Asp Gly
 1 5 10 15
 Gln Pro Val Val Leu Ala Pro Asn His Thr Asp His Cys His Tyr Gln
 20 25 30
 Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val Val Leu Cys Thr Cys
 35 40 45
 Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg Asn Ala Ser Tyr Tyr
 50 55 60
 Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp Phe Ser Thr His Glu
 65 70 75 80
 Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys Gly Thr Cys Gly His
 85 90 95
 Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser Leu Pro Gly Gly Pro
 100 105 110
 Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr Arg Lys Tyr Leu Glu
 115 120 125
 Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu Thr Arg His Arg Asn
 130 135 140

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Leu	Asn	His	Thr	Lys	Gln	Arg	Leu	Leu	Glu	Val	Ala	Asn	Tyr	Val	Asp
145					150					155					160
Gln	Leu	Leu	Arg	Thr	Leu	Asp	Ile	Gln	Val	Ala	Leu	Thr	Gly	Leu	Glu
				165					170					175	
Val	Trp	Thr	Glu	Arg	Asp	Arg	Ser	Arg	Val	Thr	Gln	Asp	Ala	Asn	Ala
			180					185					190		
Thr	Leu	Trp	Ala	Phe	Leu	Gln	Trp	Arg	Arg	Gly	Leu	Trp	Ala	Gln	Arg
		195					200					205			
Pro	His	Asp	Ser	Ala	Gln	Leu	Leu	Thr	Gly	Arg	Ala	Phe	Gln	Gly	Ala
	210					215					220				
Thr	Val	Gly	Leu	Ala	Pro	Val	Glu	Gly	Met	Cys	Arg	Ala	Glu	Ser	Ser
225					230					235					240
Gly	Gly	Val	Ser	Thr	Asp	His	Ser	Glu	Leu	Pro	Ile	Gly	Ala	Ala	Ala
				245					250					255	
Thr	Met	Ala	His	Glu	Ile	Gly	His	Ser	Leu	Gly	Leu	Ser	His	Asp	Pro
			260					265					270		
Asp	Gly	Cys	Cys	Val	Glu	Ala	Ala	Ala	Glu	Ser	Gly	Gly	Cys	Val	Met
		275					280					285			
Ala	Ala	Ala	Thr	Gly	His	Pro	Phe	Pro	Arg	Val	Phe	Ser	Ala	Cys	Ser
	290					295					300				
Arg	Arg	Gln	Leu	Arg	Ala	Phe	Phe	Arg	Lys	Gly	Gly	Gly	Ala	Cys	Leu
305					310					315					320
Ser	Asn	Ala	Pro	Asp	Pro	Gly	Leu	Pro	Val	Pro	Pro	Ala	Leu	Cys	Gly
				325					330					335	
Asn	Gly	Phe	Val	Glu	Ala	Gly	Glu	Glu	Cys	Asp	Cys	Gly	Pro	Gly	Gln
			340					345					350		
Glu	Cys	Arg	Asp	Leu	Cys	Cys	Phe	Ala	His	Asn	Cys	Ser	Leu	Arg	Pro
		355					360					365			
Gly	Ala	Gln	Cys	Ala	His	Gly	Asp	Cys	Cys	Val	Arg	Cys	Leu	Leu	Lys
	370					375					380				
Pro	Ala	Gly	Ala	Leu	Cys	Arg	Gln	Ala	Met	Gly	Asp	Cys	Asp	Leu	Pro
385					390					395					400
Glu	Phe	Cys	Thr	Gly	Thr	Ser	Ser	His	Cys	Pro	Pro	Asp	Val	Tyr	Leu
				405					410					415	
Leu	Asp	Gly	Ser	Pro	Cys	Ala	Arg	Gly	Ser	Gly	Tyr	Cys	Trp	Asp	Gly
			420					425					430		
Ala	Cys	Pro	Thr	Leu	Glu	Gln	Gln	Cys	Gln	Gln	Leu	Trp	Gly	Pro	Gly
		435					440					445			
Ser	His	Pro	Ala	Pro	Glu	Ala	Cys	Phe	Gln	Val	Val	Asn	Ser	Ala	Gly
	450					455					460				
Asp	Ala	His	Gly	Asn	Cys	Gly	Gln	Asp	Ser	Glu	Gly	His	Phe	Leu	Pro
465					470					475					480

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Cys Ala Gly Arg Asp Ala Leu Cys Gly Lys Leu Gln Cys Gln Gly Gly
 485 490 495
 Lys Pro Ser Leu Leu Ala Pro His Met Val Pro Val Asp Ser Thr Val
 500 505 510
 His Leu Asp Gly Gln Glu Val Thr Cys Arg Gly Ala Leu Ala Leu Pro
 515 520 525
 Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val Glu Pro Gly Thr
 530 535 540
 Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg Cys Arg Lys Asn
 545 550 555 560
 Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala Cys His Ser His Gly
 565 570 575
 Val Cys Asn Ser Asn His Asn Cys His Cys Ala Pro Gly Trp Ala Pro
 580 585 590
 Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met Asp Ser Gly Pro
 595 600 605
 Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu Ala Met Leu Leu Ser
 610 615 620
 Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu Ala Trp Cys Cys Tyr
 625 630 635 640
 Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser Trp Gly Cys Arg Arg
 645 650 655
 Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro His Arg Asp His Pro
 660 665 670
 Leu Gly Gly Val His Pro Met Glu Leu Gly Pro Thr Ala Thr Gly Gln
 675 680 685
 Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu Pro Ser Ser His Pro
 690 695 700
 Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro Gln Ala Asp Gln Val
 705 710 715 720
 Gln Met Pro Arg Ser Cys Leu Trp
 725

<210> 81
 <211> 802
 <212> PRT
 <213> Homo sapiens

<400> 81
 Met Gly Trp Arg Pro Arg Arg Ala Arg Gly Thr Pro Leu Leu Leu Leu
 1 5 10 15
 Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln
 20 25 30
 Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
 35 40 45

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Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
 50 55 60
 Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
 65 70 75 80
 Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
 85 90 95
 Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Asp
 100 105 110
 His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val
 115 120 125
 Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg
 130 135 140
 Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp
 145 150 155 160
 Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys
 165 170 175
 Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser
 180 185 190
 Leu Pro Gly Gly Pro Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr
 195 200 205
 Arg Lys Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu
 210 215 220
 Thr Arg His Arg Asn Leu Asn His Thr Lys Gln Arg Leu Leu Glu Val
 225 230 235 240
 Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala
 245 250 255
 Leu Thr Gly Leu Glu Val Trp Thr Glu Arg Asp Arg Ser Arg Val Thr
 260 265 270
 Gln Asp Ala Asn Ala Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg Gly
 275 280 285
 Leu Trp Ala Gln Arg Pro His Asp Ser Ala Gln Leu Leu Thr Gly Arg
 290 295 300
 Ala Phe Gln Gly Ala Thr Val Gly Leu Ala Pro Val Glu Gly Met Cys
 305 310 315 320
 Arg Ala Glu Ser Ser Gly Gly Val Ser Thr Asp His Ser Glu Leu Pro
 325 330 335
 Ile Gly Ala Ala Ala Thr Met Ala His Glu Ile Gly His Ser Leu Gly
 340 345 350
 Leu Ser His Asp Pro Asp Gly Cys Cys Val Glu Ala Ala Ala Glu Ser
 355 360 365
 Gly Gly Cys Val Met Ala Ala Ala Thr Gly His Pro Phe Pro Arg Val
 370 375 380

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Phe Ser Ala Cys Ser Arg Arg Gln Leu Arg Ala Phe Phe Arg Lys Gly
 385 390 395 400
 Gly Gly Ala Cys Leu Ser Asn Ala Pro Asp Pro Gly Leu Pro Val Pro
 405 410 415
 Pro Ala Leu Cys Gly Asn Gly Phe Val Glu Ala Gly Glu Glu Cys Asp
 420 425 430
 Cys Gly Pro Gly Gln Glu Cys Arg Asp Leu Cys Cys Phe Ala His Asn
 435 440 445
 Cys Ser Leu Arg Pro Gly Ala Gln Cys Ala His Gly Asp Cys Cys Val
 450 455 460
 Arg Cys Leu Leu Lys Pro Ala Gly Ala Leu Cys Arg Gln Ala Met Gly
 465 470 475 480
 Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly Thr Ser Ser His Cys Pro
 485 490 495
 Pro Asp Val Tyr Leu Leu Asp Gly Ser Pro Cys Ala Arg Gly Ser Gly
 500 505 510
 Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys Gln Gln
 515 520 525
 Leu Trp Gly Pro Gly Ser His Pro Ala Pro Glu Ala Cys Phe Gln Val
 530 535 540
 Val Asn Ser Ala Gly Asp Ala His Gly Asn Cys Gly Gln Asp Ser Glu
 545 550 555 560
 Gly His Phe Leu Pro Cys Ala Gly Arg Asp Ala Leu Cys Gly Lys Leu
 565 570 575
 Gln Cys Gln Gly Gly Lys Pro Ser Leu Leu Ala Pro His Met Val Pro
 580 585 590
 Val Asp Ser Thr Val His Leu Asp Gly Gln Glu Val Thr Cys Arg Gly
 595 600 605
 Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu
 610 615 620
 Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg
 625 630 635 640
 Arg Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala
 645 650 655
 Cys His Ser His Gly Ala Gly Leu His Pro Ser Val Thr Ser Gln Ala
 660 665 670
 Leu Val Ala Ala Trp Thr Val Ala Leu Cys Arg Leu Lys Thr Met Thr
 675 680 685
 Pro Ser Cys Trp Pro Cys Ser Ser Ala Ser Cys Cys Leu Cys Ser Gln
 690 695 700
 Gly Pro Ala Trp Pro Gly Val Ala Thr Asp Ser Gln Glu Pro Ile Cys
 705 710 715 720

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Ser Asp Ala Ala Gly Ala Ala Glu Gly Thr Leu Arg Ala Val Ala Pro
725 730
Lys Met Ala His Thr Gly Thr Thr Pro Trp Ala Ala Phe Thr Pro Trp
740 745 750
Ser Trp Ala Pro Gln Pro Leu Asp Ser Pro Gly Pro Trp Thr Leu Arg
755 760 765
Thr Leu Met Ser Pro Ala Ala Thr Leu Arg Ser Leu Cys Gln Gln Ser
770 775 780
Arg Leu Thr Pro Lys Ile Lys Ser Arg Cys Gln Asp Pro Ala Ser Gly
785 790 795 800
Glu Arg

<210> 82
<211> 685
<212> PRT
<213> Mus musculus

<400> 82
Asp His Cys Gln Tyr His Gly Arg Val Arg Gly Phe Arg Glu Ser Trp
1 5 10 15
Val Val Leu Ser Thr Cys Ser Gly Met Ser Gly Leu Ile Val Leu Ser
20 25 30
Ser Lys Val Ser Tyr Tyr Leu Gln Pro Arg Thr Pro Gly Asp Thr Lys
35 40 45
Asp Phe Pro Thr His Glu Ile Phe Arg Met Glu Gln Leu Phe Thr Trp
50 55 60
Arg Gly Val Gln Arg Asp Lys Asn Ser Gln Tyr Lys Ala Gly Met Ala
65 70 75 80
Ser Leu Pro His Val Pro Gln Ser Arg Val Arg Arg Glu Ala Arg Arg
85 90 95
Ser Pro Arg Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe
100 105 110
Leu Leu Gln His Gln Asn Leu Asn His Thr Arg Gln Arg Leu Leu Glu
115 120 125
Val Ala Asn Cys Val Asp Gln Ile Leu Arg Thr Leu Asp Ile Gln Leu
130 135 140
Val Leu Thr Gly Leu Glu Val Trp Thr Glu Gln Asp Leu Ser Arg Ile
145 150 155 160
Thr Gln Asp Ala Asn Glu Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg
165 170 175
Gly Val Trp Ala Arg Arg Pro His Asp Ser Thr Gln Leu Leu Thr Gly
180 185 190
Arg Thr Phe Gln Gly Thr Thr Val Gly Leu Ala Pro Val Glu Asp Met
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195	200	205
Pro Arg Gly Glu Leu Ser Phe Gly Gly Val Ser Thr Asp His Ser Glu 210 215 220		
Leu Pro Ile Gly Thr Ala Ala Thr Met Ala His Glu Ile Gly His Ser 225 230 235 240		
Leu Gly Leu His His Asp Pro Glu Gly Cys Cys Val Gln Ala Asp Ala 245 250 255		
Glu Gln Gly Gly Cys Val Met Glu Ala Ala Thr Gly His Pro Phe Pro 260 265 270		
Arg Val Phe Ser Ala Cys Ser Arg Arg Gln Leu Arg Thr Phe Phe Arg 275 280 285		
Lys Gly Gly Gly Pro Cys Leu Ser Asn Thr Ser Ala Pro Gly Leu Leu 290 295 300		
Val Leu Pro Ser Arg Cys Gly Asn Gly Phe Leu Glu Ala Gly Glu Glu 305 310 315 320		
Cys Asp Cys Gly Ser Gly Gln Lys Cys Pro Asp Pro Cys Cys Phe Ala 325 330 335		
His Asn Cys Ser Leu Arg Ala Gly Ala Gln Cys Ala His Gly Asp Cys 340 345 350		
Cys Ala Arg Cys Leu Leu Lys Ser Ala Gly Thr Pro Cys Arg Pro Ala 355 360 365		
Ala Thr Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly Thr Ser Pro Tyr 370 375 380		
Cys Pro Ala Asp Val Tyr Leu Leu Asp Gly Ser Pro Cys Ala Glu Gly 385 390 395 400		
Arg Gly Tyr Cys Leu Asp Gly Trp Cys Pro Thr Leu Glu Gln Gln Cys 405 410 415		
Gln Gln Leu Trp Gly Pro Gly Ser Lys Pro Ala Pro Glu Pro Cys Phe 420 425 430		
Gln Gln Met Asn Ser Met Gly Asn Ser Gln Gly Asn Cys Gly Gln Asp 435 440 445		
His Lys Gly Ser Phe Leu Pro Cys Ala Gln Arg Asp Ala Leu Cys Gly 450 455 460		
Lys Leu Leu Cys Gln Gly Gly Glu Pro Asn Pro Leu Val Pro His Ile 465 470 475 480		
Val Thr Met Asp Ser Thr Ile Leu Leu Glu Gly Arg Glu Val Val Cys 485 490 495		
Arg Gly Ala Phe Val Leu Pro Asp Ser His Leu Asp Gln Leu Asp Leu 500 505 510		
Gly Leu Val Glu Pro Gly Thr Gly Cys Gly Pro Arg Met Val Cys Gln 515 520 525		
Asp Arg His Cys Gln Asn Ala Thr Ser Gln Glu Leu Glu Arg Cys Leu		

530 535 540
 Thr Ala Cys His Asn Gly Gly Val Cys Asn Ser Asn Arg Asn Cys His
 545 550 555 560
 Cys Ala Ala Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Leu Gly
 565 570 575
 Gly Ser Val Asp Ser Gly Pro Ala Gln Ser Ala Asn Arg Asp Ala Phe
 580 585 590
 Pro Leu Ala Met Leu Leu Ser Phe Leu Leu Pro Leu Leu Pro Gly Ala
 595 600 605
 Gly Leu Ala Trp Cys Tyr Gln Leu Pro Thr Phe Cys His Arg Arg Gly
 610 615 620
 Leu Cys Cys Arg Arg Asp Pro Leu Trp Asn Arg Asp Ile Pro Leu Gly
 625 630 635 640
 Ser Val His Pro Val Glu Phe Gly Ser Ile Ile Thr Gly Glu Pro Ser
 645 650 655
 Pro Pro Pro Pro Trp Thr Ser Cys Gln Gln Arg Ser His Pro Pro Ser
 660 665 670
 Leu Asp Leu Leu Ser Asp Pro Ala Asn Ser Glu Leu Thr
 675 680 685

<210> 83

<211> 914

<212> PRT

<213> *Xenopus laevis*

<400> 83

Met Gly Thr Glu Gly Arg Leu Ser Thr Trp Leu Gly Leu Gly Ala Val
 1 5 10 15
 Ile Val Gly Leu Leu Leu Pro Pro Val Leu Thr Leu Gly Ala His Gln
 20 25 30
 Gly Glu Leu Val Thr Ala Phe Trp Leu Gln Asn Gly Arg Ala Lys Arg
 35 40 45
 Ser Val Asp Leu Leu Asp Lys Gly Thr Pro Asp Gly Gly Glu Ile Leu
 50 55 60
 Val Ser Ser Glu Gly Arg Lys Phe Ile Leu Lys Val Glu Arg Asn His
 65 70 75 80
 Leu Leu Phe Ala Pro Gly Tyr Thr Glu Thr His Tyr Thr Asp Gly Gln
 85 90 95
 Met Val Thr Leu Ser Pro Asn His Thr Glu His Cys Tyr Tyr His Gly
 100 105 110
 Gln Val Glu Asn Tyr Asp Glu Ser Ser Val Ala Leu Thr Thr Cys Ser
 115 120 125
 Gly Ile Ser Gly Leu Ile Trp Leu Ser Thr Asn Asn Ser Tyr Tyr Leu
 130 135 140

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Lys Pro Leu Glu Val Pro Gly Lys Glu Thr His Thr Leu Val Arg Thr
 145 150 155 160
 Glu His Leu Leu Ile Lys Glu Gly Ser Cys Gly His Asp Gly His Ser
 165 170 175
 Gly Ser Thr Ala Ser Tyr Leu Gln Glu Phe Thr Ala Pro Ser Ser His
 180 185 190
 His His Arg Val Arg Arg Asn Val Trp Arg Ser Gln Lys Tyr Met Glu
 195 200 205
 Leu Phe Ile Val Ala Asp Tyr Ser Met Phe Met Lys Gln Asn Arg Asn
 210 215 220
 Leu Gly Ser Thr Lys Gln Arg Val Leu Glu Ile Ala Asn Tyr Val Asp
 225 230 235 240
 Lys Phe Tyr Met Ser Met Asn Ile Lys Val Ala Leu Ile Gly Leu Glu
 245 250 255
 Val Trp Thr Glu Arg Asp Gln Cys Glu Val Asn Asp Asp Ala Asn Asp
 260 265 270
 Ser Leu Lys Ser Phe Leu Gln Trp Lys Gln Lys Leu Arg Ser Arg Lys
 275 280 285
 Lys His Asp Asn Ala Gln Leu Ile Thr Gly Val Thr Phe Lys Gly Thr
 290 295 300
 Thr Ile Gly Met Ala Pro Leu Glu Gly Met Cys Thr Ala Glu Asn Ser
 305 310 315 320
 Gly Gly Val Ser Met Asp His Ser Glu Asn Ala Ile Gly Ala Ala Ala
 325 330 335
 Thr Met Ala His Glu Ile Gly His Asn Phe Gly Met Ser His Asp Asp
 340 345 350
 Gly Cys Cys Val Glu Ala Thr Pro Glu Gln Gly Gly Cys Ile Met Ala
 355 360 365
 Ala Ala Thr Gly His Pro Phe Pro Arg Lys Phe Ser Ser Cys Ser Gln
 370 375 380
 Lys Gln Leu Met Ser Tyr Phe Gln Lys Gly Gly Gly Met Cys Leu Phe
 385 390 395 400
 Asn Met Pro Asn Thr Lys Asp Leu Val Met Gly Lys Lys Cys Gly Asn
 405 410 415
 Gly Phe Leu Glu Glu Gly Glu Gln Cys Asp Cys Gly Glu Pro Glu Glu
 420 425 430
 Cys Thr Asn Ser Cys Cys Asn Ala Asn Asn Cys Thr Leu Lys Ala Gly
 435 440 445
 Ala Gln Cys Ala His Gly Glu Cys Cys Gln Asp Cys Lys Leu Lys Ser
 450 455 460
 Ala Gly Thr Gln Cys Arg Glu Met Ala Gly Ser Cys Asp Leu Pro Glu
 465 470 475 480

Phe Cys Thr Gly Asp Ala Pro Ser Cys Pro Ser Asn Val Tyr Lys Leu
 485 490 495
 Asp Gly Ser Leu Cys Ala Asp Gly Asn Ala Tyr Cys Tyr Asn Gly Met
 500 505 510
 Cys Leu Thr His Gln Gln Gln Cys Ile His Leu Trp Gly Ser Gly Ala
 515 520 525
 Val Val Ala Pro Asn Phe Cys Phe Gln Asp Val Asn Lys Ala Gly Asp
 530 535 540
 Gln Tyr Gly Asn Cys Gly Lys Asn Gly Arg Gly Gln Phe Val Lys Cys
 545 550 555 560
 Thr Ser Arg Asp Ala Lys Cys Gly Lys Ile Gln Cys Gln Thr Ser Ser
 565 570 575
 Glu Lys Pro Arg Asp Pro Ser Met Val Lys Val Asp Asn Thr Ile Ile
 580 585 590
 Ile Asn Gly Tyr Lys Met Lys Cys Gln Gly Val His Ala Tyr Ser Met
 595 600 605
 Gln Glu Glu Glu Gly Asp Pro Gly Leu Val Met Thr Gly Thr Lys Cys
 610 615 620
 Gly Asp Gly Met Val Cys Lys Asp Arg Arg Cys Gln Asn Ala Ser Phe
 625 630 635 640
 Phe Glu Leu Asp Gln Cys Val Ser Lys Cys Asn Gly His Gly Val Cys
 645 650 655
 Asn Ser Asn Arg Asn Cys His Cys Asp Ser Gly Trp Ala Pro Pro Tyr
 660 665 670
 Cys Asp Lys Pro Gly Pro Gly Gly Ser Gln Asp Ser Gly Pro Ala Pro
 675 680 685
 Ser Asp Leu Pro Val Gly Val Thr Ile Phe Leu Val Ile Leu Phe Leu
 690 695 700
 Val Leu Leu Leu Ala Leu Ala Phe Ala Met Val Tyr Trp Tyr Arg Lys
 705 710 715 720
 Pro Gly Ser Leu Leu Asn Arg Trp Leu Met Lys Ser Lys Ala Lys Cys
 725 730 735
 Ser Leu Cys Lys Ala Thr Gln Pro Lys Ala Asn Arg Ala Tyr Ser Ser
 740 745 750
 Arg Ile Phe Thr Leu Arg Asn Ile Ser Tyr Pro Val Lys Ser Thr Ser
 755 760 765
 Lys Glu Thr Arg Ser Arg Asp Ile Phe Gln Gly Lys Thr Thr Ala Ala
 770 775 780
 Gln Asn Ser Ser Gln Pro Val Asn Val Val Arg Pro Leu Arg Pro Ala
 785 790 795 800
 Pro Ser Pro Val Ile Gln His Gly Val Gln Val Lys Pro Leu Arg Pro
 805 810 815

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Pro Pro Pro Pro Met Lys Pro Ser Pro Ile Leu Pro Ala Lys Glu Gln
820 825 830

Thr Val His Val Lys Leu Leu Pro Pro Lys Lys Pro Leu Pro Ser Cys
835 840 845

Pro Ile Arg Thr Gln Gln Leu Asn Pro Pro Ser Lys Pro Leu Pro Val
850 855 860

Thr Pro Ala His Lys Glu Pro Leu Leu Val Leu Thr Pro Ala Thr His
865 870 875 880

Lys Pro Pro Ile Thr Asn Ser Ala Thr Gln Leu Lys Gly Pro His Arg
885 890 895

Pro Ile Gln Gly Gly Lys Val Gln Ala Ala Ala Ala Ala Phe Leu Gln
900 905 910

Arg Lys

<210> 84
<211> 203
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Reprolysin
(M12B) family zinc metalloprotease domain sequence

<400> 84
Lys Tyr Ile Glu Leu Val Ile Val Val Asp His Gly Met Tyr Thr Lys
1 5 10 15

Tyr Gly Ser Asp Leu Asn Lys Ile Arg Gln Arg Val His Gln Ile Val
20 25 30

Asn Leu Val Asn Glu Ile Tyr Arg Pro Gln Leu Asn Ile Arg Val Val
35 40 45

Leu Val Gly Leu Glu Ile Trp Ser Asp Gly Asp Lys Ile Asn Val Gln
50 55 60

Ser Asp Ala Asn Asp Thr Leu His Ser Phe Gly Glu Trp Arg Glu Thr
65 70 75 80

Asp Leu Leu Lys Arg Lys Ser His Asp Asn Ala Gln Leu Leu Thr Gly
85 90 95

Ile Asp Phe Asp Gly Asn Thr Ile Gly Ala Ala Tyr Val Gly Gly Met
100 105 110

Cys Ser Pro Lys Arg Ser Val Gly Val Val Gln Asp His Ser Pro Ile
115 120 125

Val Leu Leu Val Ala Val Thr Met Ala His Glu Leu Gly His Asn Leu
130 135 140

Gly Met Thr His Asp Asp Lys Asn Lys Asp Gly Cys Thr Cys Glu Gly
145 150 155 160

Gly Gly Ser Cys Ile Met Asn Pro Val Ala Ser Ser Ser Pro Ser Lys

165 170 175
 Lys Lys Phe Ser Asn Cys Ser Lys Asp Asp Tyr Gln Lys Phe Leu Thr
 180 185 190
 Lys Gln Lys Pro Gln Cys Leu Leu Asn Lys Pro
 195 200

<210> 85
 <211> 119
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Pep_M12B_Propep (Reprolysin family propeptide)
 domain sequence

<400> 85
 His Leu Glu Lys Asn Arg Ser Leu Leu Ala Pro Asp Phe Thr Val Thr
 1 5 10 15
 Thr Tyr Asp Glu Asp Gly Thr Leu Val Thr Glu Glu Pro Leu Ile Gln
 20 25 30
 Asp Asp His Cys Tyr Tyr Gln Gly Tyr Val Glu Gly Tyr Pro Asn Ser
 35 40 45
 Ala Val Ser Leu Ser Thr Cys Ser Gly Gly Leu Arg Gly Ile Leu Gln
 50 55 60
 Leu Glu Asn Leu Ser Tyr Gly Ile Glu Pro Leu Glu Ser Ser Asp Gly
 65 70 75 80
 Phe Glu His Ile Ile Tyr Gln Ile Glu Asn Asp Lys Thr Glu Pro Ser
 85 90 95
 Pro Cys Gly Glu Cys Gly Ser Leu Ser Thr Ser Thr Asp Ser Ser Tyr
 100 105 110
 Gly Ile Arg Ser Ala Ser Pro
 115

<210> 86
 <211> 422
 <212> PRT
 <213> Homo sapiens

<400> 86
 Met Phe Ser Asn Ser Asp Glu Ala Val Ile Asn Lys Lys Leu Pro Lys
 1 5 10 15
 Glu Leu Leu Leu Arg Ile Phe Ser Phe Leu Asp Val Val Thr Leu Cys
 20 25 30
 Arg Cys Ala Gln Val Ser Arg Ala Trp Asn Val Leu Ala Leu Asp Gly
 35 40 45
 Ser Asn Trp Gln Arg Ile Asp Leu Phe Asp Phe Gln Arg Asp Ile Glu
 50 55 60

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Gly Arg Val Val Glu Asn Ile Ser Lys Arg Cys Gly Gly Phe Leu Arg
 65 70 75 80
 Lys Leu Ser Leu Arg Gly Cys Leu Gly Val Gly Asp Asn Ala Leu Arg
 85 90 95
 Thr Phe Ala Gln Asn Cys Arg Asn Ile Glu Val Leu Asn Leu Asn Gly
 100 105 110
 Cys Thr Lys Thr Thr Asp Ala Thr Cys Thr Ser Leu Ser Lys Phe Cys
 115 120 125
 Ser Lys Leu Arg His Leu Asp Leu Ala Ser Cys Thr Ser Ile Thr Asn
 130 135 140
 Met Ser Leu Lys Ala Leu Ser Glu Gly Cys Pro Leu Leu Glu Gln Leu
 145 150 155 160
 Asn Ile Ser Trp Cys Asp Gln Val Thr Lys Asp Gly Ile Gln Ala Leu
 165 170 175
 Val Arg Gly Cys Gly Gly Leu Lys Ala Leu Phe Leu Lys Gly Cys Thr
 180 185 190
 Gln Leu Glu Asp Glu Ala Leu Lys Tyr Ile Gly Ala His Cys Pro Glu
 195 200 205
 Leu Val Thr Leu Asn Leu Gln Thr Cys Leu Gln Ile Thr Asp Glu Gly
 210 215 220
 Leu Ile Thr Ile Cys Arg Gly Cys His Lys Leu Gln Ser Leu Cys Ala
 225 230 235 240
 Ser Gly Cys Ser Asn Ile Thr Asp Ala Ile Leu Asn Ala Leu Gly Gln
 245 250 255
 Asn Cys Pro Arg Leu Arg Ile Leu Glu Val Ala Arg Cys Ser Gln Leu
 260 265 270
 Thr Asp Val Gly Phe Thr Thr Leu Ala Arg Asn Cys His Glu Leu Glu
 275 280 285
 Lys Met Asp Leu Glu Glu Cys Val Gln Ile Thr Asp Ser Thr Leu Ile
 290 295 300
 Gln Leu Ser Ile His Cys Pro Arg Leu Gln Val Leu Ser Leu Ser His
 305 310 315 320
 Cys Glu Leu Ile Thr Asp Asp Gly Ile Arg His Leu Gly Asn Gly Ala
 325 330 335
 Cys Ala His Asp Gln Leu Glu Val Ile Glu Leu Asp Asn Cys Pro Leu
 340 345 350
 Ile Thr Asp Ala Ser Leu Glu His Leu Lys Ser Cys His Ser Leu Glu
 355 360 365
 Arg Ile Glu Leu Tyr Asp Cys Gln Gln Ile Thr Arg Ala Gly Ile Lys
 370 375 380
 Arg Leu Arg Thr His Leu Pro Asn Ile Lys Val His Ala Tyr Phe Ala
 385 390 395 400

Pro Val Thr Pro Pro Pro Ser Val Gly Gly Ser Arg Gln Arg Phe Cys
 405 410 415

Arg Cys Cys Ile Ile Leu
 420

<210> 87
 <211> 422
 <212> PRT
 <213> Mus musculus

<400> 87
 Met Phe Ser Asn Ser Asp Glu Ala Val Ile Asn Lys Lys Leu Pro Lys
 1 5 10 15
 Glu Leu Leu Leu Arg Ile Phe Ser Phe Pro Asp Val Val Thr Leu Cys
 20 25 30
 Arg Cys Ala Gln Val Ser Arg Ala Trp Asn Val Leu Ala Leu Asp Gly
 35 40 45
 Ser Asn Trp Gln Arg Ile Asp Leu Phe Asp Phe Gln Arg Asp Ile Glu
 50 55 60
 Gly Arg Val Val Glu Asn Ile Ser Lys Arg Cys Gly Gly Phe Leu Arg
 65 70 75 80
 Lys Leu Ser Leu Arg Gly Cys Leu Gly Val Gly Asp Asn Ala Leu Arg
 85 90 95
 Thr Phe Ala Gln Asn Cys Arg Asn Ile Glu Val Leu Ser Leu Asn Gly
 100 105 110
 Cys Thr Lys Thr Thr Asp Ala Thr Cys Thr Ser Leu Ser Lys Phe Cys
 115 120 125
 Ser Lys Leu Arg His Leu Asp Leu Ala Ser Cys Thr Ser Ile Thr Asn
 130 135 140
 Met Ser Leu Lys Ala Leu Ser Glu Gly Cys Pro Leu Leu Glu Gln Leu
 145 150 155 160
 Asn Ile Ser Trp Cys Asp Gln Val Thr Lys Asp Gly Ile Gln Ala Leu
 165 170 175
 Val Arg Gly Cys Gly Gly Leu Lys Ala Leu Phe Leu Lys Gly Cys Thr
 180 185 190
 Gln Leu Glu Asp Glu Ala Leu Lys Tyr Ile Gly Ala His Cys Pro Glu
 195 200 205
 Leu Val Thr Leu Asn Leu Gln Thr Cys Leu Gln Ile Thr Asp Glu Gly
 210 215 220
 Leu Ile Thr Ile Cys Arg Gly Cys His Lys Leu Gln Ser Leu Cys Ala
 225 230 235 240
 Ser Gly Cys Ser Asn Ile Thr Asp Ala Ile Leu Asn Ala Leu Gly Gln
 245 250 255
 Asn Cys Pro Arg Leu Arg Ile Leu Glu Val Ala Arg Cys Ser Gln Leu
 260 265 270

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Thr Asp Val Gly Phe Thr Thr Leu Ala Arg Asn Cys His Glu Leu Glu
 275 280 285
 Lys Met Asp Leu Glu Glu Cys Val Gln Ile Thr Asp Ser Thr Leu Ile
 290 295 300
 Gln Leu Ser Ile His Cys Pro Arg Leu Gln Val Leu Ser Leu Ser His
 305 310 315 320
 Cys Glu Leu Ile Thr Asp Asp Gly Ile Arg His Leu Gly Asn Gly Ala
 325 330 335
 Cys Ala His Asp Gln Leu Glu Val Ile Glu Leu Asp Asn Cys Pro Leu
 340 345 350
 Ile Thr Asp Ala Ser Leu Glu His Leu Lys Ser Cys Pro Ser Phe Glu
 355 360 365
 Arg Ile Glu Leu Tyr Asp Cys Gln Gln Ile Thr Arg Ala Gly Ile Lys
 370 375 380
 Arg Leu Arg Thr His Leu Pro Asn Ile Lys Val His Ala Tyr Phe Ala
 385 390 395 400
 Pro Val Thr Pro Pro Pro Ser Val Gly Gly Ser Arg Gln Arg Phe Cys
 405 410 415
 Arg Cys Cys Ile Ile Leu
 420

<210> 88
 <211> 423
 <212> PRT
 <213> Homo sapiens

<400> 88
 Met Val Phe Ser Asn Asn Asp Glu Gly Leu Ile Asn Lys Lys Leu Pro
 1 5 10 15
 Lys Glu Leu Leu Leu Arg Ile Phe Ser Phe Leu Asp Ile Val Thr Leu
 20 25 30
 Cys Arg Cys Ala Gln Ile Ser Lys Ala Trp Asn Ile Leu Ala Leu Asp
 35 40 45
 Gly Ser Asn Trp Gln Arg Ile Asp Leu Phe Asn Phe Gln Thr Asp Val
 50 55 60
 Glu Gly Arg Val Val Glu Asn Ile Ser Lys Arg Cys Gly Gly Phe Leu
 65 70 75 80
 Arg Lys Leu Ser Leu Arg Gly Cys Ile Gly Val Gly Asp Ser Ser Leu
 85 90 95
 Lys Thr Phe Ala Gln Asn Cys Arg Asn Ile Glu His Leu Asn Leu Asn
 100 105 110
 Gly Cys Thr Lys Ile Thr Asp Ser Thr Cys Tyr Ser Leu Ser Arg Phe
 115 120 125
 Cys Ser Lys Leu Lys His Leu Asp Leu Thr Ser Cys Val Ser Ile Thr
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130 135 140
 Asn Ser Ser Leu Lys Gly Ile Ser Glu Gly Cys Arg Asn Leu Glu Tyr
 145 150 155 160
 Leu Asn Leu Ser Trp Cys Asp Gln Ile Thr Lys Asp Gly Ile Glu Ala
 165 170 175
 Leu Val Arg Gly Cys Arg Gly Leu Lys Ala Leu Leu Leu Arg Gly Cys
 180 185 190
 Thr Gln Leu Glu Asp Glu Ala Leu Lys His Ile Gln Asn Tyr Cys His
 195 200 205
 Glu Leu Val Ser Leu Asn Leu Gln Ser Cys Ser Arg Ile Thr Asp Glu
 210 215 220
 Gly Val Val Gln Ile Cys Arg Gly Cys His Arg Leu Gln Ala Leu Cys
 225 230 235 240
 Leu Ser Gly Cys Ser Asn Leu Thr Asp Ala Ser Leu Thr Ala Leu Gly
 245 250 255
 Leu Asn Cys Pro Arg Leu Gln Ile Leu Glu Ala Ala Arg Cys Ser His
 260 265 270
 Leu Thr Asp Ala Gly Phe Thr Leu Leu Ala Arg Asn Cys His Glu Leu
 275 280 285
 Glu Lys Met Asp Leu Glu Glu Cys Ile Leu Ile Thr Asp Ser Thr Leu
 290 295 300
 Ile Gln Leu Ser Ile His Cys Pro Lys Leu Gln Ala Leu Ser Leu Ser
 305 310 315 320
 His Cys Glu Leu Ile Thr Asp Asp Gly Ile Leu His Leu Ser Asn Ser
 325 330 335
 Thr Cys Gly His Glu Arg Leu Arg Val Leu Glu Leu Asp Asn Cys Leu
 340 345 350
 Leu Ile Thr Asp Val Ala Leu Glu His Leu Glu Asn Cys Arg Gly Leu
 355 360 365
 Glu Arg Leu Glu Leu Tyr Asp Cys Gln Gln Val Thr Arg Ala Gly Ile
 370 375 380
 Lys Arg Met Arg Ala Gln Leu Pro His Val Lys Val His Ala Tyr Phe
 385 390 395 400
 Ala Pro Val Thr Pro Pro Thr Ala Val Ala Gly Ser Gly Gln Arg Leu
 405 410 415
 Cys Arg Cys Cys Val Ile Leu
 420

<210> 89
 <211> 425
 <212> PRT
 <213> Homo sapiens

<400> 89

Ser₁ Ala Met Val Phe₅ Ser Asn Asn Asp₁₀ Glu Gly Leu Ile Asn Lys₁₅ Lys
 Leu Pro Lys Glu₂₀ Leu Leu Leu Arg Ile₂₅ Phe Ser Phe Leu Asp₃₀ Ile Val
 Thr Leu Cys₃₅ Arg Cys Ala Gln Ile₄₀ Ser Lys Ala Trp Asn₄₅ Ile Leu Ala
 Leu Asp₅₀ Gly Ser Asn Trp Gln₅₅ Arg Ile Asp Leu Phe₆₀ Asn Phe Gln Ile
 Asp₆₅ Val Glu Gly Arg Val₇₀ Val Glu Asn Ile Ser₇₅ Lys Arg Cys Gly Gly₈₀
 Phe Leu Arg Lys₈₅ Leu Ser Leu Arg Gly Cys₉₀ Ile Gly Val Gly Asp₉₅ Ser
 Ser Leu Lys Thr₁₀₀ Phe Ala Gln Asn Cys₁₀₅ Arg Asn Ile Glu His₁₁₀ Leu Asn
 Leu Asn Gly₁₁₅ Cys Thr Lys Ile Thr₁₂₀ Asp Ser Thr Cys Tyr₁₂₅ Ser Leu Ser
 Arg Phe₁₃₀ Cys Ser Lys Leu Lys₁₃₅ His Leu Asp Leu Thr₁₄₀ Ser Cys Val Ser
 Ile Thr₁₄₅ Asn Ser Ser Leu₁₅₀ Lys Gly Ile Ser Glu₁₅₅ Gly Cys Arg Asn Leu₁₆₀
 Glu Tyr Leu Asn Leu₁₆₅ Ser Trp Cys Asp Gln Ile Thr Lys Asp Gly₁₇₅ Ile
 Glu Ala Leu Val₁₈₀ Arg Gly Cys Arg Gly₁₈₅ Leu Lys Ala Leu Leu₁₉₀ Leu Arg
 Gly Cys Thr₁₉₅ Gln Leu Glu Asp Glu₂₀₀ Ala Leu Lys His Ile₂₀₅ Gln Asn Tyr
 Cys His₂₁₀ Glu Leu Val Ser Leu₂₁₅ Asn Leu Gln Ser Cys₂₂₀ Ser Arg Ile Thr
 Asp₂₂₅ Glu Gly Val Val Gln₂₃₀ Ile Cys Arg Gly Cys₂₃₅ His Arg Leu Gln Ala₂₄₀
 Leu Cys Leu Ser Gly₂₄₅ Cys Ser Asn Leu Thr₂₅₀ Asp Ala Ser Leu Thr₂₅₅ Ala
 Leu Gly Leu Asn Cys Pro Arg Leu Gln Ile Leu Glu Ala Ala Arg Cys
 Ser His₂₇₅ Leu Thr Asp Ala Gly Phe₂₈₀ Thr Leu Leu Ala Arg₂₈₅ Asn Cys His
 Glu Leu₂₉₀ Glu Lys Met Asp Leu₂₉₅ Glu Glu Cys Ile Leu₃₀₀ Ile Thr Asp Ser
 Thr₃₀₅ Leu Ile Gln Leu Ser₃₁₀ Ile His Cys Pro Lys₃₁₅ Leu Gln Ala Leu Ser₃₂₀
 Leu Ser His Cys Glu₃₂₅ Leu Ile Thr Asp Asp₃₃₀ Gly Ile Leu His Leu Ser₃₃₅

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Asn Ser Thr Cys Gly His Glu Arg Leu Arg Val Leu Glu Leu Asp Asn
 340 345 350

Cys Leu Leu Ile Thr Asp Val Ala Leu Glu His Leu Glu Asn Cys Arg
 355 360 365

Gly Leu Glu Arg Leu Glu Leu Tyr Asp Cys Gln Gln Val Thr Arg Ala
 370 375 380

Gly Ile Lys Arg Met Arg Ala Gln Leu Pro His Val Lys Val His Ala
 385 390 395 400

Tyr Phe Ala Pro Val Thr Pro Pro Thr Ala Val Ala Gly Ser Gly Gln
 405 410 415

Arg Leu Cys Arg Cys Cys Val Ile Leu
 420 425

<210> 90
 <211> 423
 <212> PRT
 <213> Homo sapiens

<400> 90
 Met Val Phe Ser Asn Asn Asp Glu Gly Leu Ile Asn Lys Lys Leu Pro
 1 5 10 15

Lys Glu Leu Leu Leu Arg Ile Phe Ser Phe Leu Asp Ile Val Thr Leu
 20 25 30

Cys Arg Cys Ala Gln Ile Ser Lys Ala Trp Asn Ile Leu Ala Leu Asp
 35 40 45

Gly Ser Asn Trp Gln Arg Ile Asp Leu Phe Asn Phe Gln Thr Asp Val
 50 55 60

Glu Gly Arg Val Val Glu Asn Ile Ser Lys Arg Cys Gly Gly Phe Leu
 65 70 75 80

Lys Lys Leu Ser Leu Arg Gly Cys Ile Gly Val Gly Asp Ser Ser Leu
 85 90 95

Lys Thr Phe Ala Gln Asn Cys Arg Asn Ile Glu His Leu Asn Leu Asn
 100 105 110

Gly Cys Thr Lys Ile Thr Asp Ser Thr Cys Tyr Ser Leu Ser Arg Phe
 115 120 125

Cys Ser Lys Leu Lys His Leu Asp Leu Thr Ser Cys Val Ser Ile Thr
 130 135 140

Asn Ser Ser Leu Lys Gly Ile Ser Glu Gly Cys Arg Asn Leu Glu Tyr
 145 150 155 160

Leu Asn Leu Ser Trp Cys Asp Gln Ile Thr Lys Asp Gly Ile Glu Ala
 165 170 175

Leu Val Arg Gly Cys Arg Gly Leu Lys Ala Leu Leu Leu Arg Gly Cys
 180 185 190

Thr Gln Leu Glu Asp Glu Ala Leu Lys His Ile Gln Asn Tyr Cys His
 195 200 205

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Glu Leu Val Ser Leu Asn Leu Gln Ser Cys Ser Arg Ile Thr Asp Glu
 210 215 220
 Gly Val Val Gln Ile Cys Arg Gly Cys His Arg Leu Gln Ala Leu Cys
 225 230 235 240
 Leu Ser Gly Cys Ser Asn Leu Thr Asp Ala Ser Leu Thr Ala Leu Gly
 245 250 255
 Leu Asn Cys Pro Arg Leu Gln Ile Leu Glu Ala Ala Arg Cys Ser His
 260 265 270
 Leu Thr Asp Ala Gly Phe Thr Leu Leu Ala Arg Asn Cys His Glu Leu
 275 280 285
 Glu Lys Met Asp Leu Glu Glu Cys Ile Leu Ile Thr Asp Ser Thr Leu
 290 295 300
 Ile Gln Leu Ser Ile His Cys Pro Lys Leu Gln Ala Leu Ser Leu Ser
 305 310 315 320
 His Cys Glu Leu Ile Thr Asp Asp Gly Ile Leu His Leu Ser Asn Ser
 325 330 335
 Thr Cys Gly His Glu Arg Leu Arg Val Leu Glu Leu Asp Asn Cys Leu
 340 345 350
 Leu Ile Thr Asp Val Ala Leu Glu His Leu Glu Asn Cys Arg Gly Leu
 355 360 365
 Glu Arg Leu Glu Leu Tyr Asp Cys Gln Gln Val Thr Arg Ala Gly Ile
 370 375 380
 Lys Arg Met Arg Ala Gln Leu Pro His Val Lys Val His Ala Tyr Phe
 385 390 395 400
 Ala Pro Val Thr Pro Pro Thr Ala Val Ala Gly Ser Gly Gln Arg Leu
 405 410 415
 Cys Arg Cys Cys Val Ile Leu
 420

<210> 91
 <211> 46
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: F-box domain
 sequence

<400> 91
 Phe Ser Leu Leu Arg Leu Pro Asp Asp Leu Leu Glu Lys Ile Leu Ser
 1 5 10 15
 Arg Leu Pro Leu Lys Asp Leu Leu Ser Leu Ser Lys Val Ser Lys Lys
 20 25 30
 Phe Arg Ser Leu Val Asp Ser Leu Leu Asp Val Lys Leu Leu
 35 40 45

<210> 92
 <211> 172
 <212> PRT
 <213> Homo sapiens

<400> 92
 Met Val Gly Pro Ala Pro Arg Arg Arg Leu Arg Pro Leu Ala Ala Leu
 1 5 10 15
 Ala Leu Val Leu Ala Leu Ala Pro Gly Leu Pro Thr Ala Arg Ala Gly
 20 25 30
 Gln Thr Pro Arg Pro Ala Glu Arg Gly Pro Pro Val Arg Leu Phe Thr
 35 40 45
 Glu Glu Glu Leu Ala Arg Tyr Gly Gly Glu Glu Glu Asp Gln Pro Ile
 50 55 60
 Tyr Leu Ala Val Lys Gly Val Val Phe Asp Val Thr Ser Gly Lys Glu
 65 70 75 80
 Phe Tyr Gly Arg Gly Ala Pro Tyr Asn Ala Leu Thr Gly Lys Asp Ser
 85 90 95
 Thr Arg Gly Val Ala Lys Met Ser Leu Asp Pro Ala Asp Leu Thr His
 100 105 110
 Asp Thr Thr Gly Leu Thr Ala Lys Glu Leu Glu Ala Leu Asp Glu Val
 115 120 125
 Phe Thr Lys Val Tyr Lys Ala Lys Tyr Pro Ile Val Gly Tyr Thr Ala
 130 135 140
 Arg Arg Ile Leu Asn Glu Asp Gly Ser Pro Asn Leu Asp Phe Lys Pro
 145 150 155 160
 Glu Asp Gln Pro His Phe Asp Ile Lys Asp Glu Phe
 165 170

<210> 93
 <211> 171
 <212> PRT
 <213> Mus musculus

<400> 93
 Met Ala Arg Pro Ala Pro Trp Trp Arg Leu Arg Leu Leu Ala Ala Leu
 1 5 10 15
 Val Leu Ala Leu Ala Leu Val Pro Val Pro Ser Ala Trp Ala Gly Gln
 20 25 30
 Thr Pro Arg Pro Ala Glu Arg Gly Pro Pro Val Arg Leu Phe Thr Glu
 35 40 45
 Glu Glu Leu Ala Arg Tyr Gly Gly Glu Glu Glu Asp Gln Pro Ile Tyr
 50 55 60
 Leu Ala Val Lys Gly Val Val Phe Asp Val Thr Ser Gly Lys Glu Phe
 65 70 75 80
 Tyr Gly Arg Gly Ala Pro Tyr Asn Ala Leu Ala Gly Lys Asp Ser Ser
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85 90 95
 Arg Gly Val Ala Lys Met Ser Leu Asp Pro Ala Asp Leu Thr His Asp
 100 105 110
 Thr Thr Gly Leu Thr Ala Lys Glu Leu Glu Ala Leu Asp Asp Val Phe
 115 120 125
 Ser Lys Val Tyr Lys Ala Lys Tyr Pro Ile Val Gly Tyr Thr Ala Arg
 130 135 140
 Arg Ile Leu Asn Glu Asp Gly Ser Pro Asn Leu Asp Phe Lys Pro Glu
 145 150 155 160
 Asp Gln Pro His Phe Asp Ile Lys Asp Glu Phe
 165 170

<210> 94
 <211> 100
 <212> PRT
 <213> Arabidopsis thaliana

<400> 94
 Met Glu Phe Thr Ala Glu Gln Leu Ser Gln Tyr Asn Gly Thr Asp Glu
 1 5 10 15
 Ser Lys Pro Ile Tyr Val Ala Ile Lys Gly Arg Val Phe Asp Val Thr
 20 25 30
 Thr Gly Lys Ser Phe Tyr Gly Ser Gly Gly Asp Tyr Ser Met Phe Ala
 35 40 45
 Gly Lys Asp Ala Ser Arg Ala Leu Gly Lys Met Ser Lys Asn Glu Glu
 50 55 60
 Asp Val Ser Pro Ser Leu Glu Gly Leu Thr Glu Lys Glu Ile Asn Thr
 65 70 75 80
 Leu Asn Asp Trp Glu Thr Lys Phe Glu Ala Lys Tyr Pro Val Val Gly
 85 90 95
 Arg Val Val Ser
 100

<210> 95
 <211> 232
 <212> PRT
 <213> Oryza sativa

<400> 95
 Met Ala Ala Ala Val Ala Glu Leu Trp Glu Thr Leu Lys Gln Ala Ile
 1 5 10 15
 Val Ala Tyr Thr Gly Leu Ser Pro Ala Ala Phe Phe Thr Ala Val Ala
 20 25 30
 Ala Ala Ala Ala Leu Tyr His Val Val Ser Gly Ile Phe Ala Gly Pro
 35 40 45
 Pro Pro Pro Pro Pro Arg Pro Arg Asp Glu Pro Glu Ala Glu Pro
 50 55 60

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Leu Pro Pro Pro Val Gln Leu Gly Glu Val Ser Glu Glu Glu Leu Arg
 65 70 75 80
 Gln Tyr Asp Gly Ser Asp Pro Lys Lys Pro Leu Leu Met Ala Ile Lys
 85 90 95
 Gly Gln Ile Tyr Asp Val Thr Gln Ser Arg Met Phe Tyr Gly Pro Gly
 100 105 110
 Gly Pro Tyr Ala Leu Phe Ala Gly Lys Asp Ala Ser Arg Ala Leu Ala
 115 120 125
 Lys Met Ser Phe Glu Pro Gln Asp Leu Thr Gly Asp Ile Ser Gly Leu
 130 135 140
 Gly Pro Phe Glu Leu Asp Ala Leu Gln Asp Trp Glu Tyr Lys Phe Met
 145 150 155 160
 Gly Lys Tyr Val Lys Val Gly Thr Val Lys Lys Thr Val Pro Val Glu
 165 170 175
 Asp Gly Ala Pro Ser Thr Ser Pro Glu Thr Thr Glu Thr Ala Ala Ala
 180 185 190
 Ala Glu Pro Glu Lys Ala Pro Ala Thr Glu Glu Lys Pro Arg Glu Val
 195 200 205
 Ser Ser Glu Glu Val Lys Glu Lys Glu Asp Ala Val Ala Ala Ala Ala
 210 215 220
 Pro Asp Glu Gly Ala Lys Glu Ser
 225 230

<210> 96

<211> 104

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Steroid
binding domain sequence

<400> 96

Asp Phe Thr Pro Glu Glu Leu Arg Lys Tyr Asp Gly Ser Asp Glu Asp
 1 5 10 15
 Lys Pro Ile Tyr Leu Ala Ile Lys Gly Lys Val Tyr Asp Val Thr Arg
 20 25 30
 Gly Arg Lys Phe Tyr Gly Pro Gly Gly Pro Tyr Ser Leu Phe Ala Gly
 35 40 45
 Arg Asp Ala Ser Arg Ala Leu Ala Thr Met Ser Phe Asp Glu Glu Asp
 50 55 60
 Leu Lys Asp Ser Asp Glu Glu Ile Asp Asp Leu Ser Asp Leu Ser Ala
 65 70 75 80
 Asp Glu Leu Glu Ala Leu Arg Glu Trp Glu Thr Lys Phe Lys Ala Lys
 85 90 95

Tyr Pro Val Val Gly Arg Leu Ile
100

<210> 97
<211> 309
<212> PRT
<213> Homo sapiens

<400> 97
Met Glu Ala Leu Ala Leu Val Gly Ala Trp Tyr Thr Ala Arg Lys Ser
1 5 10 15
Ile Thr Val Ile Cys Asp Phe Tyr Ser Leu Ile Arg Leu His Phe Ile
20 25 30
Pro Arg Leu Gly Ser Arg Ala Asp Leu Ile Lys Gln Tyr Gly Arg Trp
35 40 45
Ala Val Val Ser Gly Ala Thr Asp Gly Ile Gly Lys Ala Tyr Ala Glu
50 55 60
Glu Leu Ala Ser Arg Gly Leu Asn Ile Ile Leu Ile Ser Arg Asn Glu
65 70 75 80
Glu Lys Leu Gln Val Val Ala Lys Asp Ile Ala Asp Thr Tyr Lys Val
85 90 95
Glu Thr Asp Ile Ile Val Ala Asp Phe Ser Ser Gly Arg Glu Ile Tyr
100 105 110
Leu Pro Ile Arg Glu Ala Leu Lys Asp Lys Asp Val Gly Ile Leu Val
115 120 125
Asn Asn Val Gly Val Phe Tyr Pro Tyr Pro Gln Tyr Phe Thr Gln Leu
130 135 140
Ser Glu Asp Lys Leu Trp Asp Ile Ile Asn Val Asn Ile Ala Ala Ala
145 150 155 160
Ser Leu Met Val His Val Val Leu Pro Gly Met Val Glu Arg Lys Lys
165 170 175
Gly Ala Ile Val Thr Ile Ser Ser Gly Ser Cys Cys Lys Pro Thr Pro
180 185 190
Gln Leu Ala Ala Phe Ser Ala Ser Lys Ala Tyr Leu Asp His Phe Ser
195 200 205
Arg Ala Leu Gln Tyr Glu Tyr Ala Ser Lys Gly Ile Phe Val Gln Ser
210 215 220
Leu Ile Pro Phe Tyr Val Ala Thr Ser Met Thr Ala Pro Ser Asn Phe
225 230 235 240
Leu His Arg Cys Ser Trp Leu Val Pro Ser Pro Lys Val Tyr Ala His
245 250 255
His Ala Val Ser Thr Leu Gly Ile Ser Lys Arg Thr Thr Gly Tyr Trp
260 265 270
Ser His Ser Ile Gln Phe Leu Phe Ala Gln Tyr Met Pro Glu Trp Leu
275 280 285

Trp Val Trp Gly Ala Asn Ile Leu Asn Arg Ser Leu Arg Lys Glu Ala
 290 295 300

Leu Ser Cys Thr Ala
 305

<210> 98
 <211> 339
 <212> PRT
 <213> Drosophila melanogaster

<400> 98
 Met Gln Pro Val Leu Glu Val Ser Ile Tyr Thr Leu Leu Lys Met Ala
 1 5 10 15

Phe Ile Trp Gln Leu Ile Ser Ala Ala Ile Tyr Leu Val Gly Leu Leu
 20 25 30

Thr Ile Gly Val Phe Leu Tyr Asp Asn Leu Lys Ser Leu Val Ser Ile
 35 40 45

Ile Lys Ala Val Leu Glu Pro Tyr Phe Gln Pro His Leu Pro Arg Thr
 50 55 60

Leu Val Asp Lys Phe Gly Gln Trp Ala Val Val Thr Gly Ala Thr Asp
 65 70 75 80

Gly Ile Gly Lys Glu Tyr Ala Arg Glu Leu Ala Arg Gln Gly Ile Asn
 85 90 95

Leu Val Leu Ile Ser Arg Thr Lys Glu Lys Leu Ile Ala Val Thr Asn
 100 105 110

Glu Ile Glu Ser Gln Tyr Lys Val Lys Thr Lys Trp Ile Ala Ala Asp
 115 120 125

Phe Ala Lys Gly Arg Glu Val Tyr Asp Gln Ile Glu Lys Glu Leu Ala
 130 135 140

Gly Ile Asp Val Gly Ile Leu Val Asn Asn Val Gly Met Met Tyr Glu
 145 150 155 160

His Pro Glu Ser Leu Asp Leu Val Ser Glu Asp Leu Leu Trp Asn Leu
 165 170 175

Leu Thr Val Asn Met Gly Ser Val Thr Met Leu Thr Arg Lys Ile Leu
 180 185 190

Pro Gln Met Ile Gly Arg Arg Lys Gly Ala Ile Val Asn Leu Gly Ser
 195 200 205

Ser Ser Glu Leu Gln Pro Leu Pro Asn Met Thr Val Tyr Ala Ala Ser
 210 215 220

Lys Lys Phe Val Thr Tyr Phe Ser Lys Ala Leu Glu Leu Glu Val Ala
 225 230 235 240

Glu His Asn Ile His Val Gln Leu Val Met Pro Asn Phe Val Val Thr
 245 250 255

Lys Met Asn Ala Tyr Thr Asp Arg Val Met Gln Gly Gly Leu Phe Phe
 Page 225

260 265 270
 Pro Asn Ala Tyr Thr Phe Ala Arg Ser Ala Val Phe Thr Leu Gly Lys
 275 280 285
 Thr Ser Glu Thr Asn Gly Phe Trp Thr His Gly Ile Gln Tyr Ala Ile
 290 295 300
 Met Lys Leu Ala Pro Leu Pro Ile Arg Thr Tyr Leu Gly His Gln Leu
 305 310 315 320
 Phe Lys Arg Leu Arg Ile Glu Ala Leu Glu Gln Lys Gln Lys Lys Leu
 325 330 335
 Lys Leu Thr

<210> 99
 <211> 312
 <212> PRT
 <213> Homo sapiens

<400> 99
 Met Glu Ser Ala Leu Pro Ala Ala Gly Phe Leu Tyr Trp Val Gly Ala
 1 5 10 15
 Gly Thr Val Ala Tyr Leu Ala Leu Arg Ile Ser Tyr Ser Leu Phe Thr
 20 25 30
 Ala Leu Arg Val Trp Gly Val Gly Asn Glu Ala Gly Val Gly Pro Gly
 35 40 45
 Leu Gly Glu Trp Ala Val Val Thr Gly Ser Thr Asp Gly Ile Gly Lys
 50 55 60
 Ser Tyr Ala Glu Glu Leu Ala Lys His Gly Met Lys Val Val Leu Ile
 65 70 75 80
 Ser Arg Ser Lys Asp Lys Leu Asp Gln Val Ser Ser Glu Ile Lys Glu
 85 90 95
 Lys Phe Lys Val Glu Thr Arg Thr Ile Ala Val Asp Phe Ala Ser Glu
 100 105 110
 Asp Ile Tyr Asp Lys Ile Lys Thr Gly Leu Ala Gly Leu Glu Ile Gly
 115 120 125
 Ile Leu Val Asn Asn Val Gly Met Ser Tyr Glu Tyr Pro Glu Tyr Phe
 130 135 140
 Leu Asp Val Pro Asp Leu Asp Asn Val Ile Lys Lys Met Ile Asn Ile
 145 150 155 160
 Asn Ile Leu Ser Val Cys Lys Met Thr Gln Leu Val Leu Pro Gly Met
 165 170 175
 Val Glu Arg Ser Lys Gly Ala Ile Leu Asn Ile Ser Ser Gly Ser Gly
 180 185 190
 Met Leu Pro Val Pro Leu Leu Thr Ile Tyr Ser Ala Thr Lys Thr Phe
 195 200 205

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Val Asp Phe Phe Ser Gln Cys Leu His Glu Glu Tyr Arg Ser Lys Gly
 210 215 220

Val Phe Val Gln Ser Val Leu Pro Tyr Phe Val Ala Thr Lys Leu Ala
 225 230 235 240

Lys Ile Arg Lys Pro Thr Leu Asp Lys Pro Ser Pro Glu Thr Phe Val
 245 250 255

Lys Ser Ala Ile Lys Thr Val Gly Leu Gln Ser Arg Thr Asn Gly Tyr
 260 265 270

Leu Ile His Ala Leu Met Gly Ser Ile Ile Ser Asn Leu Pro Ser Trp
 275 280 285

Ile Tyr Leu Lys Ile Val Met Asn Met Asn Lys Ser Thr Arg Ala His
 290 295 300

Tyr Leu Lys Lys Thr Lys Lys Asn
 305 310

<210> 100
 <211> 312
 <212> PRT
 <213> Anas platyrhynchos

<400> 100
 Met Leu Pro Ala Ala Gly Leu Leu Trp Trp Val Gly Ala Leu Gly Ala
 1 5 10 15

Leu Tyr Ala Ala Val Arg Gly Ala Leu Gly Leu Leu Gly Ala Leu Arg
 20 25 30

Val Trp Gly Ile Gly Ala Gly Arg Ala Ala Leu Gly Pro Gly Leu Gly
 35 40 45

Ala Trp Ala Val Val Thr Gly Ala Thr Asp Gly Ile Gly Lys Ala Tyr
 50 55 60

Ala Lys Glu Leu Ala Lys Arg Gly Met Lys Val Ala Leu Ile Ser Arg
 65 70 75 80

Ser Lys Glu Lys Leu Asp Gln Val Ala Gly Glu Ile Thr Glu Gln Tyr
 85 90 95

Gly Val Glu Thr Lys Val Ile Val Ala Asp Phe Gly Glu Arg Glu Asp
 100 105 110

Ile Tyr Asp Arg Ile Arg Ala Gly Leu Glu Gly Leu Glu Ile Gly Val
 115 120 125

Leu Val Asn Asn Val Gly Ile Ser Tyr Ser Tyr Pro Glu Tyr Phe Ile
 130 135 140

Asp Val Pro Asp Leu Asp Lys Thr Ile Asp Lys Met Ile Asn Ile Asn
 145 150 155 160

Ile Met Ser Val Cys Lys Met Thr Arg Leu Val Leu Pro Gly Met Leu
 165 170 175

Glu Arg Ser Lys Gly Val Ile Leu Asn Ile Ser Ser Ala Ala Gly Met
 180 185 190

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Tyr Pro Thr Pro Leu Leu Thr Leu Tyr Ser Ala Ser Lys Ala Phe Val
 195 200 205
 Asp Tyr Phe Ser Arg Gly Leu His Ala Glu Tyr Lys Ser Lys Gly Ile
 210 215 220
 Ile Val Gln Ser Val Met Pro Tyr Tyr Val Ala Thr Lys Met Ser Lys
 225 230 235 240
 Ile Ser Lys Pro Ser Phe Asp Lys Pro Thr Pro Glu Thr Tyr Val Arg
 245 250 255
 Ala Ala Ile Gly Thr Val Gly Leu Gln Ser Gln Thr Asn Gly Cys Leu
 260 265 270
 Pro His Ala Phe Met Gly Trp Val Phe Ser Ile Leu Pro Thr Ser Thr
 275 280 285
 Val Met Asn Leu Leu Met Lys Thr Asn Lys Gln Ile Arg Ala Arg Phe
 290 295 300
 Leu Lys Lys Lys Met Lys Glu Lys
 305 310

<210> 101
 <211> 312
 <212> PRT
 <213> Mus musculus

<400> 101
 Met Glu Cys Ala Pro Pro Ala Ala Gly Phe Leu Tyr Trp Val Gly Ala
 1 5 10 15
 Ser Thr Ile Ala Tyr Leu Ala Leu Arg Ala Ser Tyr Ser Leu Phe Arg
 20 25 30
 Ala Phe Gln Val Trp Cys Val Gly Asn Glu Ala Leu Val Gly Pro Arg
 35 40 45
 Leu Gly Glu Trp Ala Val Val Thr Gly Gly Thr Asp Gly Ile Gly Lys
 50 55 60
 Ala Tyr Ala Glu Glu Leu Ala Lys Arg Gly Met Lys Ile Val Leu Ile
 65 70 75 80
 Ser Arg Ser Gln Asp Lys Leu Asn Gln Val Ser Asn Asn Ile Lys Glu
 85 90 95
 Lys Phe Asn Val Glu Thr Arg Thr Ile Ala Val Asp Phe Ser Leu Asp
 100 105 110
 Asp Ile Tyr Asp Lys Ile Lys Thr Gly Leu Ser Gly Leu Glu Ile Gly
 115 120 125
 Val Leu Val Asn Asn Val Gly Met Ser Tyr Glu Tyr Pro Glu Tyr Phe
 130 135 140
 Leu Glu Ile Pro Asp Leu Asp Asn Thr Ile Lys Lys Leu Ile Asn Ile
 145 150 155 160
 Asn Val Leu Ser Val Cys Lys Val Thr Arg Leu Val Leu Pro Gly Met
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165 170 175
 Val Glu Arg Ser Lys Gly Val Ile Leu Asn Ile Ser Ser Ala Ser Gly
 180 185 190
 Met Leu Pro Val Pro Leu Leu Thr Ile Tyr Ser Ala Thr Lys Ala Phe
 195 200 205
 Val Asp Phe Phe Ser Gln Cys Leu His Glu Glu Tyr Lys Ser Lys Gly
 210 215 220
 Ile Phe Val Gln Ser Val Met Pro Tyr Leu Val Ala Thr Lys Leu Ala
 225 230 235 240
 Lys Ile Gln Lys Pro Thr Leu Asp Lys Pro Ser Ala Glu Thr Phe Val
 245 250 255
 Lys Ser Ala Ile Lys Thr Val Gly Leu Gln Thr Arg Thr Thr Gly Tyr
 260 265 270
 Val Ile His Ser Leu Met Gly Ser Ile Asn Ser Ile Met Pro Arg Trp
 275 280 285
 Met Tyr Phe Lys Ile Ile Met Gly Phe Ser Lys Ser Leu Arg Asn Arg
 290 295 300
 Tyr Leu Lys Lys Arg Lys Lys Asn
 305 310

<210> 102

<211> 271

<212> PRT

<213> Artificial Sequence

<220>

 <223> Description of Artificial Sequence: Short Chain
 Alcohol Dehydrogenase (adh_short) domain sequence

<400> 102

Thr Gly Lys Val Ala Leu Val Thr Gly Ala Ser Ser Gly Ile Gly Leu
 1 5 10 15
 Ala Ile Ala Lys Arg Leu Ala Lys Glu Gly Ala Lys Val Val Val Val
 20 25 30
 Asp Arg Arg Glu Glu Lys Ala Glu Gln Val Ala Ala Glu Leu Lys Ala
 35 40 45
 Glu Leu Gly Asp Arg Ala Leu Phe Ile Gln Leu Asp Val Thr Asp Glu
 50 55 60
 Glu Gln Val Lys Ala Ala Val Ala Gln Ala Val Glu Arg Leu Gly Asp
 65 70 75 80
 Arg Leu Asp Val Leu Val Asn Asn Ala Gly Ile Leu Gly Pro Gly Pro
 85 90 95
 Pro Phe Glu Glu Leu Ser Glu Glu Asp Trp Glu Arg Val Ile Asp Val
 100 105 110
 Asn Leu Thr Gly Val Phe Leu Leu Thr Gln Ala Val Leu Pro Ala Met
 115 120 125

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Asp His Met Leu Lys Arg Lys Gly Gly Arg Ile Val Asn Ile Ser Ser
 130 135 140
 Val Ala Gly Leu Asn Val Gly Val Pro Gly Leu Ser Ala Tyr Ser Ala
 145 150 155 160
 Ser Lys Ala Ala Val Ile Gly Leu Thr Arg Ser Leu Ala Leu Glu Leu
 165 170 175
 Ala Pro His Gly Thr Gly Ile Arg Val Asn Ala Val Ala Pro Gly Gly
 180 185 190
 Val Asp Thr Asp Met Thr Lys Ala Leu Arg Ser Arg Leu Ile Glu Ala
 195 200 205
 Lys Lys Lys Val Arg Glu Val Ala Asp Ile Ala Asp Pro Glu Leu Glu
 210 215 220
 Glu Arg Ile Thr Ser Thr Ile Thr Pro Leu Gly Arg Tyr Gly Val Thr
 225 230 235 240
 Pro Glu Glu Ile Ala Asn Ala Val Leu Phe Leu Ala Ser Asp Gly Ala
 245 250 255
 Ser Tyr Ser Val Thr Gly Gln Thr Leu Asn Val Asp Gly Gly Leu
 260 265 270

<210> 103
 <211> 1961
 <212> PRT
 <213> Homo sapiens

<400> 103
 Met Ala Gln Gln Ala Ala Asp Lys Tyr Leu Tyr Val Asp Lys Asn Phe
 1 5 10 15
 Ile Asn Asn Pro Leu Ala Gln Ala Asp Trp Ala Ala Lys Lys Leu Val
 20 25 30
 Trp Val Pro Ser Asp Lys Ser Gly Phe Glu Pro Ala Ser Leu Lys Glu
 35 40 45
 Glu Val Gly Glu Arg Gly His Val Glu Leu Val Glu Asn Gly Lys Lys
 50 55 60
 Val Lys Val Asn Lys Asp Asp Ile Gln Lys Met Asn Pro Pro Lys Phe
 65 70 75 80
 Ser Lys Val Glu Asp Met Ala Glu Leu Thr Cys Leu Asn Glu Ala Ser
 85 90 95
 Val Leu His Asn Leu Lys Glu Arg Tyr Tyr Ser Gly Leu Ile Tyr Thr
 100 105 110
 Tyr Ser Gly Leu Phe Cys Val Val Ile Asn Pro Tyr Lys Asn Leu Pro
 115 120 125
 Ile Tyr Ser Glu Glu Ile Val Glu Met Tyr Lys Gly Lys Lys Arg His
 130 135 140
 Glu Met Pro Pro His Ile Tyr Ala Ile Thr Asp Thr Ala Tyr Arg Ser
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145					150						155					160
Met	Met	Gln	Asp	Arg	Glu	Asp	Gln	Ser	Ile	Leu	Cys	Thr	Gly	Glu	Ser	
				165					170					175		
Gly	Ala	Gly	Lys	Thr	Glu	Asn	Thr	Lys	Lys	Val	Ile	Gln	Tyr	Leu	Ala	
			180					185					190			
Tyr	Val	Ala	Ser	Ser	His	Lys	Ser	Lys	Lys	Asp	Gln	Gly	Glu	Leu	Glu	
		195					200					205				
Arg	Gln	Leu	Leu	Gln	Ala	Asn	Pro	Ile	Leu	Glu	Ala	Phe	Gly	Asn	Ala	
	210					215					220					
Lys	Thr	Val	Lys	Asn	Asp	Asn	Ser	Ser	Arg	Phe	Gly	Lys	Phe	Ile	Arg	
225					230					235					240	
Ile	Asn	Phe	Asp	Val	Asn	Gly	Tyr	Ile	Val	Gly	Ala	Asn	Ile	Glu	Thr	
				245					250					255		
Tyr	Leu	Leu	Glu	Lys	Ser	Arg	Ala	Ile	Arg	Gln	Ala	Lys	Glu	Glu	Arg	
			260					265					270			
Thr	Phe	His	Ile	Phe	Tyr	Tyr	Leu	Leu	Ser	Gly	Ala	Gly	Glu	His	Leu	
		275					280					285				
Lys	Thr	Asp	Leu	Leu	Leu	Glu	Pro	Tyr	Asn	Lys	Tyr	Arg	Phe	Leu	Ser	
	290					295					300					
Asn	Gly	His	Val	Thr	Ile	Pro	Gly	Gln	Gln	Asp	Lys	Asp	Met	Phe	Gln	
305					310					315					320	
Glu	Thr	Met	Glu	Ala	Met	Arg	Ile	Met	Gly	Ile	Pro	Glu	Glu	Glu	Gln	
				325					330					335		
Met	Gly	Leu	Leu	Arg	Val	Ile	Ser	Gly	Val	Leu	Gln	Leu	Gly	Asn	Ile	
			340					345					350			
Val	Phe	Lys	Lys	Glu	Arg	Asn	Thr	Asp	Gln	Ala	Ser	Met	Pro	Asp	Asn	
		355					360					365				
Thr	Ala	Ala	Gln	Lys	Val	Ser	His	Leu	Leu	Gly	Ile	Asn	Val	Thr	Asp	
	370					375					380					
Phe	Thr	Arg	Gly	Ile	Leu	Thr	Pro	Arg	Ile	Lys	Val	Gly	Arg	Asp	Tyr	
385					390					395					400	
Val	Gln	Lys	Ala	Gln	Thr	Lys	Glu	Gln	Ala	Asp	Phe	Ala	Ile	Glu	Ala	
				405					410					415		
Leu	Ala	Lys	Ala	Thr	Tyr	Glu	Arg	Met	Phe	Arg	Trp	Leu	Val	Leu	Arg	
			420					425					430			
Ile	Asn	Lys	Ala	Leu	Asp	Lys	Thr	Lys	Arg	Gln	Gly	Ala	Ser	Phe	Ile	
		435					440					445				
Gly	Ile	Leu	Asp	Ile	Ala	Gly	Phe	Glu	Ile	Phe	Asp	Leu	Asn	Ser	Phe	
	450					455					460					
Glu	Gln	Leu	Cys	Ile	Asn	Tyr	Thr	Asn	Glu	Lys	Leu	Gln	Gln	Leu	Phe	
465					470					475					480	
Asn	His	Thr	Met	Phe	Ile	Leu	Glu	Gln	Glu	Glu	Tyr	Gln	Arg	Glu	Gly	

820				825				830						
Lys Val	Lys 835	Pro	Leu	Leu	Gln	Val 840	Ser	Arg	Gln	Glu	Glu 845	Glu	Met	Met
Ala	Lys 850	Glu	Glu	Glu	Leu	Val 855	Lys	Val	Arg	Glu	Lys 860	Gln	Leu	Ala
Glu 865	Asn	Arg	Leu	Met	Glu 870	Met	Glu	Thr	Leu	Gln 875	Ser	Gln	Leu	Met
Glu	Lys	Leu	Gln	Leu 885	Gln	Glu	Gln	Leu	Gln 890	Ala	Glu	Thr	Glu	Leu
Ala	Glu	Ala	Glu 900	Glu	Leu	Arg	Ala	Arg 905	Leu	Thr	Ala	Lys	Lys 910	Gln
Leu	Glu	Glu 915	Ile	Cys	His	Asp	Leu 920	Glu	Ala	Arg	Val	Glu 925	Glu	Glu
Glu	Arg 930	Tyr	Gln	His	Leu	Gln 935	Ala	Glu	Lys	Lys	Lys 940	Met	Gln	Gln
Ile 945	Gln	Glu	Leu	Glu	Glu 950	Gln	Leu	Glu	Glu	Glu 955	Glu	Ser	Ala	Arg
Lys	Leu	Gln	Leu	Glu 965	Lys	Val	Thr	Thr	Glu 970	Ala	Lys	Leu	Lys	Lys 975
Glu	Glu	Glu	Gln 980	Ile	Ile	Leu	Glu	Asp 985	Gln	Asn	Cys	Lys	Leu	Ala
Glu	Lys	Lys 995	Leu	Leu	Glu	Asp	Arg 1000	Ile	Ala	Glu	Phe	Thr 1005	Thr	Asn
Thr	Glu 1010	Glu	Glu	Glu	Lys	Ser 1015	Lys	Ser	Leu	Ala	Lys 1020	Leu	Lys	Asn
His 1025	Glu	Ala	Met	Ile	Thr 1030	Asp	Leu	Glu	Glu	Arg 1035	Leu	Arg	Arg	Glu
Lys	Gln	Arg	Gln 1045	Glu	Leu	Glu	Lys	Thr	Arg 1050	Arg	Lys	Leu	Glu	Gly 1055
Ser	Thr	Asp	Leu 1060	Ser	Asp	Gln	Ile 1065	Ala	Glu	Leu	Gln	Ala	Gln 1070	Ile
Glu	Leu	Lys 1075	Met	Gln	Leu	Ala	Lys 1080	Lys	Glu	Glu	Glu	Leu	Gln 1085	Ala
Leu	Ala	Arg	Val 1090	Glu	Glu	Glu 1095	Ala	Ala	Gln	Lys	Asn 1100	Met	Ala	Leu
Lys	Ile 1105	Arg	Glu	Leu	Glu 1110	Ser	Gln	Ile	Ser	Glu 1115	Leu	Gln	Glu	Asp
Glu	Ser	Glu	Arg	Ala 1125	Ser	Arg	Asn	Lys	Ala 1130	Glu	Lys	Gln	Lys	Arg
Leu	Gly	Glu	Glu 1140	Leu	Glu	Ala	Leu	Lys 1145	Thr	Glu	Leu	Glu	Asp 1150	Thr
Asp	Ser	Thr	Ala	Ala	Gln	Gln	Glu	Leu	Arg	Ser	Lys	Arg	Glu	Gln

1155 1160 1165
 Val Asn Ile Leu Lys Lys Thr Leu Glu Glu Glu Ala Lys Thr His Glu
 1170 1175 1180
 Ala Gln Ile Gln Glu Met Arg Gln Lys His Ser Gln Ala Val Glu Glu
 1185 1190 1195 1200
 Leu Ala Glu Gln Leu Glu Gln Thr Lys Arg Val Lys Ala Asn Leu Glu
 1205 1210 1215
 Lys Ala Lys Gln Thr Leu Glu Asn Glu Arg Gly Glu Leu Ala Asn Glu
 1220 1225 1230
 Val Lys Val Leu Leu Gln Gly Gly Arg Asp Ser Glu His Lys Arg Lys
 1235 1240 1245
 Lys Val Glu Ala Gln Leu Gln Glu Leu Gln Val Lys Phe Asn Glu Gly
 1250 1255 1260
 Glu Arg Val Arg Thr Glu Leu Ala Asp Lys Val Thr Lys Leu Gln Val
 1265 1270 1275 1280
 Glu Leu Asp Asn Val Thr Gly Leu Leu Ser Gln Ser Asp Ser Lys Ser
 1285 1290 1295
 Ser Lys Leu Thr Lys Asp Phe Ser Ala Leu Glu Ser Gln Leu Gln Asp
 1300 1305 1310
 Thr Gln Glu Leu Leu Gln Glu Glu Asn Arg Gln Lys Leu Ser Leu Ser
 1315 1320 1325
 Thr Lys Leu Lys Gln Val Glu Asp Glu Lys Asn Ser Phe Arg Glu Gln
 1330 1335 1340
 Leu Glu Glu Glu Glu Glu Glu Ala Lys His Asn Leu Glu Lys Gln Ile
 1345 1350 1355 1360
 Ala Thr Leu His Ala Gln Val Ala Asp Met Lys Lys Lys Met Glu Asp
 1365 1370 1375
 Ser Val Gly Cys Leu Glu Thr Ala Glu Glu Val Lys Arg Lys Leu Gln
 1380 1385 1390
 Lys Asp Leu Glu Gly Leu Ser Gln Arg His Glu Glu Lys Val Ala Ala
 1395 1400 1405
 Tyr Asp Lys Leu Glu Lys Thr Lys Thr Arg Leu Gln Gln Glu Leu Asp
 1410 1415 1420
 Asp Leu Leu Val Asp Leu Asp His Gln Arg Gln Ser Ala Cys Asn Leu
 1425 1430 1435 1440
 Glu Lys Lys Gln Lys Lys Phe Asp Gln Leu Leu Ala Glu Glu Lys Thr
 1445 1450 1455
 Ile Ser Ala Lys Tyr Ala Glu Glu Arg Asp Arg Ala Glu Ala Glu Ala
 1460 1465 1470
 Arg Glu Lys Glu Thr Lys Ala Leu Ser Leu Ala Arg Ala Leu Glu Glu
 1475 1480 1485
 Ala Met Glu Gln Lys Ala Glu Leu Glu Arg Leu Asn Lys Gln Phe Arg
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1490 1495 1500
 Thr Glu Met Glu Asp Leu Met Ser Ser Lys Asp Asp Val Gly Lys Ser
 1505 1510 1515 1520
 Val His Glu Leu Glu Lys Ser Lys Arg Ala Leu Glu Gln Gln Val Glu
 1525 1530 1535
 Glu Met Lys Thr Gln Leu Glu Glu Leu Glu Asp Glu Leu Gln Ala Thr
 1540 1545 1550
 Glu Asp Ala Lys Leu Arg Leu Glu Val Asn Leu Gln Ala Met Lys Ala
 1555 1560 1565
 Gln Phe Glu Arg Asp Leu Gln Gly Arg Asp Glu Gln Ser Glu Glu Lys
 1570 1575 1580
 Lys Lys Gln Leu Val Arg Gln Val Arg Glu Met Glu Ala Glu Leu Glu
 1585 1590 1595 1600
 Asp Glu Arg Lys Gln Arg Ser Met Ala Val Ala Ala Arg Lys Lys Leu
 1605 1610 1615
 Glu Met Asp Leu Lys Asp Leu Glu Ala His Ile Asp Ser Ala Asn Lys
 1620 1625 1630
 Asn Arg Asp Glu Ala Ile Lys Gln Leu Arg Lys Leu Gln Ala Gln Met
 1635 1640 1645
 Lys Asp Cys Met Arg Glu Leu Asp Asp Thr Arg Ala Ser Arg Glu Glu
 1650 1655 1660
 Ile Leu Ala Gln Ala Lys Glu Asn Glu Lys Lys Leu Lys Ser Met Glu
 1665 1670 1675 1680
 Ala Glu Met Ile Gln Leu Gln Glu Glu Leu Ala Ala Ala Glu Arg Ala
 1685 1690 1695
 Lys Arg Gln Ala Gln Gln Glu Arg Asp Glu Leu Ala Asp Glu Ile Ala
 1700 1705 1710
 Asn Ser Ser Gly Lys Gly Ala Leu Ala Leu Glu Glu Lys Arg Arg Leu
 1715 1720 1725
 Glu Ala Arg Ile Ala Gln Leu Glu Glu Glu Leu Glu Glu Glu Gln Gly
 1730 1735 1740
 Asn Thr Glu Leu Ile Asn Asp Arg Leu Lys Lys Ala Asn Leu Gln Ile
 1745 1750 1755 1760
 Asp Gln Ile Asn Ala Asp Leu Asn Leu Glu Arg Gly His Ala Gln Lys
 1765 1770 1775
 Asn Glu Asn Ala Arg Gln Gln Leu Glu Arg Gln Asn Lys Glu Leu Lys
 1780 1785 1790
 Val Lys Leu Gln Glu Met Glu Gly Thr Val Lys Ser Lys Tyr Lys Ala
 1795 1800 1805
 Ser Ile Thr Ala Leu Glu Ala Lys Ile Ala Gln Leu Glu Glu Gln Leu
 1810 1815 1820
 Asp Asn Glu Thr Lys Glu Arg Gln Ala Ala Cys Lys Gln Val Arg Arg
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1825	1830	1835	1840
Thr Glu Lys Lys	Leu Lys Asp Val	Leu Leu Gln Val	Asp Asp Glu Arg
	1845	1850	1855
Arg Asn Ala Glu	Gln Tyr Lys Asp	Gln Ala Asp Lys	Ala Ser Thr Arg
	1860	1865	1870
Leu Lys Gln Leu	Lys Arg Gln Leu	Glu Glu Ala Glu	Glu Glu Ala Gln
	1875	1880	1885
Arg Ala Asn Ala	Ser Arg Arg Lys	Leu Gln Arg Glu	Leu Glu Asp Ala
	1890	1895	1900
Thr Glu Thr Ala	Asp Ala Met Asn	Arg Glu Val Ser	Ser Leu Lys Asn
	1910	1915	1920
Lys Leu Arg Arg	Gly Asp Leu Pro	Phe Val Val Pro	Arg Arg Met Ala
	1925	1930	1935
Arg Lys Gly Ala	Gly Asp Gly Ser	Asp Glu Glu Val	Asp Gly Lys Ala
	1940	1945	1950
Asp Gly Ala Glu	Ala Lys Pro Ala	Glu	
	1955	1960	

<210> 104
 <211> 1960
 <212> PRT
 <213> Homo sapiens

<400> 104

Met	Ala	Gln	Gln	Ala	Ala	Asp	Lys	Tyr	Leu	Tyr	Val	Asp	Lys	Asn	Phe
1				5					10					15	
Ile	Asn	Asn	Pro	Leu	Ala	Gln	Ala	Asp	Trp	Ala	Ala	Lys	Lys	Leu	Val
			20					25					30		
Trp	Val	Pro	Ser	Asp	Lys	Ser	Gly	Phe	Glu	Pro	Ala	Ser	Leu	Lys	Glu
		35					40					45			
Glu	Val	Gly	Glu	Glu	Ala	Ile	Val	Glu	Leu	Val	Glu	Asn	Gly	Lys	Lys
	50					55					60				
Val	Lys	Val	Asn	Lys	Asp	Asp	Ile	Gln	Lys	Met	Asn	Pro	Pro	Lys	Phe
	65				70					75					80
Ser	Lys	Val	Glu	Asp	Met	Ala	Glu	Leu	Thr	Cys	Leu	Asn	Glu	Ala	Ser
				85					90					95	
Val	Leu	His	Asn	Leu	Lys	Glu	Arg	Tyr	Tyr	Ser	Gly	Leu	Ile	Tyr	Thr
			100					105					110		
Tyr	Ser	Gly	Leu	Phe	Cys	Val	Val	Ile	Asn	Pro	Tyr	Lys	Asn	Leu	Pro
		115					120					125			
Ile	Tyr	Ser	Glu	Glu	Ile	Val	Glu	Met	Tyr	Lys	Gly	Lys	Lys	Arg	His
	130					135					140				
Glu	Met	Pro	Pro	His	Ile	Tyr	Ala	Ile	Thr	Asp	Thr	Ala	Tyr	Arg	Ser
	145				150					155					160

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Met	Met	Gln	Asp	Arg 165	Glu	Asp	Gln	Ser	Ile 170	Leu	Cys	Thr	Gly	Glu 175	Ser
Gly	Ala	Gly	Lys 180	Thr	Glu	Asn	Thr	Lys 185	Lys	Val	Ile	Gln	Tyr 190	Leu	Ala
Tyr	Val	Ala 195	Ser	Ser	His	Lys	Ser 200	Lys	Lys	Asp	Gln	Gly 205	Glu	Leu	Glu
Arg	Gln 210	Leu	Leu	Gln	Ala	Asn 215	Pro	Ile	Leu	Glu	Ala 220	Phe	Gly	Asn	Ala
Lys 225	Thr	Val	Lys	Asn	Asp 230	Asn	Ser	Ser	Arg	Phe 235	Gly	Lys	Phe	Ile	Arg 240
Ile	Asn	Phe	Asp	Val 245	Asn	Gly	Tyr	Ile	Val 250	Gly	Ala	Asn	Ile	Glu 255	Thr
Tyr	Leu	Leu	Glu 260	Lys	Ser	Arg	Ala	Ile 265	Arg	Gln	Ala	Lys	Glu 270	Glu	Arg
Thr	Phe	His 275	Ile	Phe	Tyr	Tyr	Leu 280	Leu	Ser	Gly	Ala	Gly 285	Glu	His	Leu
Lys	Thr 290	Asp	Leu	Leu	Leu	Glu 295	Pro	Tyr	Asn	Lys	Tyr 300	Arg	Phe	Leu	Ser
Asn 305	Gly	His	Val	Thr	Ile 310	Pro	Gly	Gln	Gln	Asp 315	Lys	Asp	Met	Phe	Gln 320
Glu	Thr	Met	Glu	Ala 325	Met	Arg	Ile	Met	Gly 330	Ile	Pro	Glu	Glu	Glu 335	Gln
Met	Gly	Leu	Leu 340	Arg	Val	Ile	Ser	Gly 345	Val	Leu	Gln	Leu	Gly 350	Asn	Ile
Val	Phe	Lys 355	Lys	Glu	Arg	Asn	Thr 360	Asp	Gln	Ala	Ser	Met 365	Pro	Asp	Asn
Thr	Ala 370	Ala	Gln	Lys	Val	Ser 375	His	Leu	Leu	Gly	Ile 380	Asn	Val	Thr	Asp
Phe 385	Thr	Arg	Gly	Ile	Leu 390	Thr	Pro	Arg	Ile	Lys 395	Val	Gly	Arg	Asp	Tyr 400
Val	Gln	Lys	Ala	Gln 405	Thr	Lys	Glu	Gln	Ala 410	Asp	Phe	Ala	Ile	Glu 415	Ala
Leu	Ala	Lys	Ala 420	Thr	Tyr	Glu	Arg	Met	Phe	Arg	Trp	Leu	Val 430	Leu	Arg
Ile	Asn	Lys 435	Ala	Leu	Asp	Lys	Thr 440	Lys	Arg	Gln	Gly	Ala 445	Ser	Phe	Ile
Gly	Ile 450	Leu	Asp	Ile	Ala	Gly 455	Phe	Glu	Ile	Phe	Asp 460	Leu	Asn	Ser	Phe
Glu 465	Gln	Leu	Cys	Ile	Asn 470	Tyr	Thr	Asn	Glu	Lys 475	Leu	Gln	Gln	Leu	Phe 480
Asn	His	Thr	Met	Phe 485	Ile	Leu	Glu	Gln	Glu 490	Glu	Tyr	Gln	Arg	Glu 495	Gly

Ile Glu Trp Asn Phe Ile Asp Phe Gly Leu Asp Leu Gln Pro Cys Ile
 500 505 510
 Asp Leu Ile Glu Lys Pro Ala Gly Pro Pro Gly Ile Leu Ala Leu Leu
 515 520 525
 Asp Glu Glu Cys Trp Phe Pro Lys Ala Thr Asp Lys Ser Phe Val Glu
 530 535 540
 Lys Val Met Gln Glu Gln Gly Thr His Pro Lys Phe Gln Lys Pro Lys
 545 550 555 560
 Gln Leu Lys Asp Lys Ala Asp Phe Cys Ile Ile His Tyr Ala Gly Lys
 565 570 575
 Val Asp Tyr Lys Ala Asp Glu Trp Leu Met Lys Asn Met Asp Pro Leu
 580 585 590
 Asn Asp Asn Ile Ala Thr Leu Leu His Gln Ser Ser Asp Lys Phe Val
 595 600 605
 Ser Glu Leu Trp Lys Asp Val Asp Arg Ile Ile Gly Leu Asp Gln Val
 610 615 620
 Ala Gly Met Ser Glu Thr Ala Leu Pro Gly Ala Phe Lys Thr Arg Lys
 625 630 635 640
 Gly Met Phe Arg Thr Val Gly Gln Leu Tyr Lys Glu Gln Leu Ala Lys
 645 650 655
 Leu Met Ala Thr Leu Arg Asn Thr Asn Pro Asn Phe Val Arg Cys Ile
 660 665 670
 Ile Pro Asn His Glu Lys Lys Ala Gly Lys Leu Asp Pro His Leu Val
 675 680 685
 Leu Asp Gln Leu Arg Cys Asn Gly Val Leu Glu Gly Ile Arg Ile Cys
 690 695 700
 Arg Gln Gly Phe Pro Asn Arg Val Val Phe Gln Glu Phe Arg Gln Arg
 705 710 715 720
 Tyr Glu Ile Leu Thr Pro Asn Ser Ile Pro Lys Gly Phe Met Asp Gly
 725 730 735
 Lys Gln Ala Cys Val Leu Met Ile Lys Ala Leu Glu Leu Asp Ser Asn
 740 745 750
 Leu Tyr Arg Ile Gly Gln Ser Lys Val Phe Phe Arg Ala Gly Val Leu
 755 760 765
 Ala His Leu Glu Glu Glu Arg Asp Leu Lys Ile Thr Asp Val Ile Ile
 770 775 780
 Gly Phe Gln Ala Cys Cys Arg Gly Tyr Leu Ala Arg Lys Ala Phe Ala
 785 790 795 800
 Lys Arg Gln Gln Gln Leu Thr Ala Met Lys Val Leu Gln Arg Asn Cys
 805 810 815
 Ala Ala Tyr Leu Lys Leu Arg Asn Trp Gln Trp Trp Arg Leu Phe Thr
 820 825 830

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Lys Val	Lys 835	Pro	Leu	Leu	Gln	Val 840	Ser	Arg	Gln	Glu 845	Glu	Met	Met
Ala	Lys 850	Glu	Glu	Glu	Leu	Val 855	Lys	Val	Arg	Glu	Lys 860	Gln	Leu
Glu 865	Asn	Arg	Leu	Thr	Glu 870	Met	Glu	Thr	Leu	Gln 875	Ser	Gln	Leu
Glu	Lys	Leu	Gln 885	Leu	Gln	Glu	Gln	Leu	Gln 890	Ala	Glu	Thr	Glu
Ala	Glu	Ala	Glu 900	Glu	Leu	Arg	Ala	Arg 905	Leu	Thr	Ala	Lys	Lys 910
Leu	Glu	Glu 915	Ile	Cys	His	Asp	Leu 920	Glu	Ala	Arg	Val	Glu 925	Glu
Glu	Arg 930	Cys	Gln	His	Leu	Gln 935	Ala	Glu	Lys	Lys	Lys 940	Met	Gln
Ile 945	Gln	Glu	Leu	Glu	Glu 950	Gln	Leu	Glu	Glu	Glu 955	Glu	Ser	Ala
Lys	Leu	Gln	Leu	Glu 965	Lys	Val	Thr	Thr	Glu 970	Ala	Lys	Leu	Lys
Glu	Glu	Glu	Gln 980	Ile	Ile	Leu	Glu	Asp 985	Gln	Asn	Cys	Lys	Leu
Glu	Lys	Lys 995	Leu	Leu	Glu	Asp	Arg 1000	Ile	Ala	Glu	Phe	Thr 1005	Thr
Thr 1010	Glu	Glu	Glu	Glu	Lys	Ser	Lys 1015	Ser	Leu	Ala	Lys	Leu	Lys
His 1025	Glu	Ala	Met	Ile	Thr 1030	Asp	Leu	Glu	Glu	Arg 1035	Leu	Arg	Arg
Lys	Gln	Arg	Gln	Glu 1045	Leu	Glu	Lys	Thr	Arg 1050	Arg	Lys	Leu	Glu
Ser	Thr	Asp	Leu 1060	Ser	Asp	Gln	Ile	Ala 1065	Glu	Leu	Gln	Ala	Gln
Glu	Leu	Lys 1075	Met	Gln	Leu	Ala	Lys 1080	Lys	Glu	Glu	Glu	Leu 1085	Gln
Leu	Ala	Arg	Val	Glu	Glu	Glu	Ala 1095	Ala	Gln	Lys	Asn	Met	Ala
Lys 1105	Ile	Arg	Glu	Leu	Glu 1110	Ser	Gln	Ile	Ser	Glu 1115	Leu	Gln	Glu
Glu	Ser	Glu	Arg	Ala 1125	Ser	Arg	Asn	Lys	Ala 1130	Glu	Lys	Gln	Lys
Leu	Gly	Glu	Glu 1140	Leu	Glu	Ala	Leu	Lys 1145	Thr	Glu	Leu	Glu	Asp
Asp	Ser	Thr 1155	Ala	Ala	Gln	Gln	Glu 1160	Leu	Arg	Ser	Lys	Arg 1165	Glu

Val Asn Ile Leu Lys Lys Thr Leu Glu Glu Glu Ala Lys Thr His Glu
 1170 1175 1180
 Ala Gln Ile Gln Glu Met Arg Gln Lys His Ser Gln Ala Val Glu Glu
 1185 1190 1195 1200
 Leu Ala Glu Gln Leu Glu Gln Thr Lys Arg Val Lys Ala Asn Leu Glu
 1205 1210 1215
 Lys Ala Lys Gln Thr Leu Glu Asn Glu Arg Gly Glu Leu Ala Asn Glu
 1220 1225 1230
 Val Lys Val Leu Leu Gln Gly Lys Gly Asp Ser Glu His Lys Arg Lys
 1235 1240 1245
 Lys Val Glu Ala Gln Leu Gln Glu Leu Gln Val Lys Phe Asn Glu Gly
 1250 1255 1260
 Glu Arg Val Arg Thr Glu Leu Ala Asp Lys Val Thr Lys Leu Gln Val
 1265 1270 1275 1280
 Glu Leu Asp Asn Val Thr Gly Leu Leu Ser Gln Ser Asp Ser Lys Ser
 1285 1290 1295
 Ser Lys Leu Thr Lys Asp Phe Ser Ala Leu Glu Ser Gln Leu Gln Asp
 1300 1305 1310
 Thr Gln Glu Leu Leu Gln Glu Glu Asn Arg Gln Lys Leu Ser Leu Ser
 1315 1320 1325
 Thr Lys Leu Lys Gln Val Glu Asp Glu Lys Asn Ser Phe Arg Glu Gln
 1330 1335 1340
 Leu Glu Glu Glu Glu Glu Ala Lys His Asn Leu Glu Lys Gln Ile Ala
 1345 1350 1355 1360
 Thr Leu His Ala Gln Val Ala Asp Met Lys Lys Lys Met Glu Asp Ser
 1365 1370 1375
 Val Gly Cys Leu Glu Thr Ala Glu Glu Val Lys Arg Lys Leu Gln Lys
 1380 1385 1390
 Asp Leu Glu Gly Leu Ser Gln Arg His Glu Glu Lys Val Ala Ala Tyr
 1395 1400 1405
 Asp Lys Leu Glu Lys Thr Lys Thr Arg Leu Gln Gln Glu Leu Asp Asp
 1410 1415 1420
 Leu Leu Val Asp Leu Asp His Gln Arg Gln Ser Ala Cys Asn Leu Glu
 1425 1430 1435 1440
 Lys Lys Gln Lys Lys Phe Asp Gln Leu Leu Ala Glu Glu Lys Thr Ile
 1445 1450 1455
 Ser Ala Lys Tyr Ala Glu Glu Arg Asp Arg Ala Glu Ala Glu Ala Arg
 1460 1465 1470
 Glu Lys Glu Thr Lys Ala Leu Ser Leu Ala Arg Ala Leu Glu Glu Ala
 1475 1480 1485
 Met Glu Gln Lys Ala Glu Leu Glu Arg Leu Asn Lys Gln Phe Arg Thr
 1490 1495 1500

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Glu Met Glu Asp Leu Met Ser Ser Lys Asp Asp Val Gly Lys Ser Val
 1505 1510 1515 1520
 His Glu Leu Glu Lys Ser Lys Arg Ala Leu Glu Gln Gln Val Glu Glu
 1525 1530 1535
 Met Lys Thr Gln Leu Glu Glu Leu Glu Asp Glu Leu Gln Ala Thr Glu
 1540 1545 1550
 Asp Ala Lys Leu Arg Leu Glu Val Asn Leu Gln Ala Met Lys Ala Gln
 1555 1560 1565
 Phe Glu Arg Asp Leu Gln Gly Arg Asp Glu Gln Ser Glu Glu Lys Lys
 1570 1575 1580
 Lys Gln Leu Val Arg Gln Val Arg Glu Met Glu Ala Glu Leu Glu Asp
 1585 1590 1595 1600
 Glu Arg Lys Gln Arg Ser Met Ala Val Ala Ala Arg Lys Lys Leu Glu
 1605 1610 1615
 Met Asp Leu Lys Asp Leu Glu Ala His Ile Asp Ser Ala Asn Lys Asn
 1620 1625 1630
 Arg Asp Glu Ala Ile Lys Gln Leu Arg Lys Leu Gln Ala Gln Met Lys
 1635 1640 1645
 Asp Cys Met Arg Glu Leu Asp Asp Thr Arg Ala Ser Arg Glu Glu Ile
 1650 1655 1660
 Leu Ala Gln Ala Lys Glu Asn Glu Lys Lys Leu Lys Ser Met Glu Ala
 1665 1670 1675 1680
 Glu Met Ile Gln Leu Gln Glu Glu Leu Ala Ala Ala Glu Arg Ala Lys
 1685 1690 1695
 Arg Gln Ala Gln Gln Glu Arg Asp Glu Leu Ala Asp Glu Ile Ala Asn
 1700 1705 1710
 Ser Ser Gly Lys Gly Ala Leu Ala Leu Glu Glu Lys Arg Arg Leu Glu
 1715 1720 1725
 Ala Arg Ile Ala Gln Leu Glu Glu Glu Leu Glu Glu Gln Gly Asn
 1730 1735 1740
 Thr Glu Leu Ile Asn Asp Arg Leu Lys Lys Ala Asn Leu Gln Ile Asp
 1745 1750 1755 1760
 Gln Ile Asn Thr Asp Leu Asn Leu Glu Arg Ser His Ala Gln Lys Asn
 1765 1770 1775
 Glu Asn Ala Arg Gln Gln Leu Glu Arg Gln Asn Lys Glu Leu Lys Val
 1780 1785 1790
 Lys Leu Gln Glu Met Glu Gly Thr Val Lys Ser Lys Tyr Lys Ala Ser
 1795 1800 1805
 Ile Thr Ala Leu Glu Ala Lys Ile Ala Gln Leu Glu Glu Gln Leu Asp
 1810 1815 1820
 Asn Glu Thr Lys Glu Arg Gln Ala Ala Cys Lys Gln Val Arg Arg Thr
 1825 1830 1835 1840

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Glu Lys Lys Leu Lys Asp Val Leu Leu Gln Val Asp Asp Glu Arg Arg
 1845 1850 1855
 Asn Ala Glu Gln Tyr Lys Asp Gln Ala Asp Lys Ala Ser Thr Arg Leu
 1860 1865 1870
 Lys Gln Leu Lys Arg Gln Leu Glu Glu Ala Glu Glu Glu Ala Gln Arg
 1875 1880 1885
 Ala Asn Ala Ser Arg Arg Lys Leu Gln Arg Glu Leu Glu Asp Ala Thr
 1890 1895 1900
 Glu Thr Ala Asp Ala Met Asn Arg Glu Val Ser Ser Leu Lys Asn Lys
 1905 1910 1915 1920
 Leu Arg Arg Gly Asp Leu Pro Phe Val Val Pro Arg Arg Met Ala Arg
 1925 1930 1935
 Lys Gly Ala Gly Asp Gly Ser Asp Glu Glu Val Asp Gly Lys Ala Asp
 1940 1945 1950
 Gly Ala Glu Ala Lys Pro Ala Glu
 1955 1960

<210> 105
 <211> 1961
 <212> PRT
 <213> Rattus norvegicus

<400> 105
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 Ile Asn Asn Pro Leu Ala Gln Ala Asp Cys Gly Ala Lys Lys Leu Val
 20 25 30
 Trp Val Pro Ser Thr Lys Asn Gly Phe Glu Pro Ala Ser Leu Lys Glu
 35 40 45
 Glu Val Gly Glu Glu Ala Ile Val Glu Leu Val Glu Asn Gly Lys Lys
 50 55 60
 Val Lys Val Asn Lys Asp Asp Ile Gln Lys Met Asn Pro Pro Lys Phe
 65 70 75 80
 Ser Lys Val Glu Asp Met Ala Glu Leu Thr Cys Leu Asn Glu Ala Ser
 85 90 95
 Val Leu His Asn Leu Lys Glu Arg Tyr Tyr Ser Gly Leu Ile Tyr Thr
 100 105 110
 Tyr Ser Gly Leu Phe Cys Val Val Ile Asn Pro Tyr Lys Asn Leu Pro
 115 120 125
 Ile Tyr Ser Glu Glu Ile Val Asp Met Tyr Lys Gly Lys Lys Arg His
 130 135 140
 Glu Met Pro Pro His Ile Tyr Ala Ile Thr Asp Thr Ala Tyr Arg Ser
 145 150 155 160
 Met Met Gln Asp Arg Glu Asp Gln Ser Ile Leu Cys Thr Gly Glu Ser
 165 170 175

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Gly Ala Gly Lys Thr Glu Asn Thr Lys Lys Val Ile Gln Tyr Leu Ala
 180 185 190
 His Val Ala Ser Ser His Lys Ser Lys Lys Asp Gln Gly Glu Leu Glu
 195 200 205
 Arg Gln Leu Leu Gln Ala Asn Pro Ile Leu Glu Ala Phe Gly Asn Ala
 210 215 220
 Lys Thr Val Lys Asn Asp Asn Ser Ser Arg Phe Gly Lys Phe Ile Arg
 225 230 235 240
 Ile Asn Phe Asp Val Asn Gly Tyr Ile Val Gly Ala Asn Ile Glu Thr
 245 250 255
 Tyr Leu Leu Glu Lys Ser Arg Ala Ile Arg Gln Ala Lys Glu Glu Arg
 260 265 270
 Thr Phe His Ile Phe Tyr Tyr Leu Leu Ser Gly Ala Gly Glu His Leu
 275 280 285
 Lys Thr Asp Leu Leu Leu Glu Pro Tyr Asn Lys Tyr Arg Phe Leu Ser
 290 295 300
 Asn Gly His Val Thr Ile Pro Gly Gln Gln Asp Lys Asp Met Phe Gln
 305 310 315 320
 Glu Thr Met Glu Ala Met Arg Ile Met Gly Ile Pro Glu Asp Glu Gln
 325 330 335
 Met Gly Leu Leu Arg Val Ile Ser Gly Val Leu Gln Leu Gly Asn Ile
 340 345 350
 Val Phe Lys Lys Glu Arg Asn Thr Asp Gln Ala Ser Met Pro Asp Asn
 355 360 365
 Thr Ala Ala Gln Lys Val Ser His Leu Leu Gly Ile Asn Val Thr Asp
 370 375 380
 Phe Thr Arg Gly Ile Leu Thr Pro Arg Ile Lys Val Gly Arg Asp Tyr
 385 390 395 400
 Val Gln Lys Ala Gln Thr Lys Glu Gln Ala Asp Phe Ala Ile Glu Ala
 405 410 415
 Leu Ala Lys Ala Thr Tyr Glu Arg Met Phe Arg Trp Leu Val Leu Arg
 420 425 430
 Ile Asn Lys Ala Leu Asp Lys Thr Lys Arg Gln Gly Ala Ser Phe Ile
 435 440 445
 Gly Ile Leu Asp Ile Ala Gly Phe Glu Ile Phe Asp Leu Asn Ser Phe
 450 455 460
 Glu Gln Leu Cys Ile Asn Tyr Thr Asn Glu Lys Leu Gln Gln Leu Phe
 465 470 475 480
 Asn His Thr Met Phe Ile Leu Glu Gln Glu Glu Tyr Gln Arg Glu Gly
 485 490 495
 Ile Glu Trp Asn Phe Ile Asp Phe Gly Leu Asp Leu Gln Pro Cys Ile
 500 505 510

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Asp Leu Ile Glu Lys Pro Ala Gly Pro Pro Gly Ile Leu Ala Leu Leu
 515 520 525
 Asp Glu Glu Cys Trp Phe Pro Lys Ala Thr Asp Lys Ser Phe Val Glu
 530 535 540
 Lys Val Val Gln Glu Gln Gly Thr His Pro Lys Phe Gln Lys Pro Lys
 545 550 555 560
 Gln Leu Lys Asp Lys Ala Asp Phe Cys Ile Ile His Tyr Ala Gly Lys
 565 570 575
 Val Asp Tyr Lys Ala Asp Glu Trp Leu Met Lys Asn Met Asp Pro Leu
 580 585 590
 Asn Asp Asn Ile Ala Thr Leu Leu His Gln Ser Ser Asp Lys Phe Val
 595 600 605
 Ser Glu Leu Trp Lys Asp Val Asp Arg Ile Ile Gly Leu Asp Gln Val
 610 615 620
 Ala Gly Met Ser Glu Thr Ala Leu Pro Gly Ala Phe Lys Thr Arg Lys
 625 630 635 640
 Gly Met Phe Arg Thr Val Gly Gln Leu Tyr Lys Glu Gln Leu Ala Lys
 645 650 655
 Leu Met Ala Thr Leu Arg Asn Thr Asn Pro Asn Phe Val Cys Cys Ile
 660 665 670
 Ile Pro Asn His Glu Lys Lys Ala Gly Lys Leu Asp Pro His Leu Val
 675 680 685
 Leu Asp Gln Leu Arg Cys Asn Gly Val Leu Glu Gly Ile Arg Ile Cys
 690 695 700
 Arg Gln Gly Phe Pro Asn Arg Val Val Phe Gln Glu Phe Arg Gln Arg
 705 710 715 720
 Tyr Glu Ile Leu Thr Pro Asn Ser Ile Pro Lys Gly Phe Met Asp Gly
 725 730 735
 Lys Gln Ala Cys Val Leu Met Ile Lys Ala Leu Glu Leu Asp Ser Asn
 740 745 750
 Leu Tyr Arg Ile Gly Gln Ser Lys Val Phe Phe Arg Ser Gly Val Leu
 755 760 765
 Ala His Leu Glu Glu Glu Arg Asp Leu Lys Ile Thr Asp Val Ile Ile
 770 775 780
 Gly Phe Gln Ala Cys Cys Arg Gly Tyr Leu Ala Arg Lys Ala Phe Ala
 785 790 795 800
 Lys Arg Gln Gln Gln Leu Thr Ala Met Lys Val Leu Gln Arg Asn Cys
 805 810 815
 Ala Ala Tyr Leu Arg Leu Arg Asn Trp Gln Trp Trp Arg Leu Phe Thr
 820 825 830
 Lys Val Lys Pro Leu Leu Asn Ser Ile Arg His Glu Asp Glu Leu Leu
 835 840 845

Ala Lys Glu Ala Glu Leu Thr Lys Val Arg Glu Lys His Leu Ala Ala
 850 855 860
 Glu Asn Arg Leu Thr Glu Met Glu Thr Met Gln Ser Gln Leu Met Ala
 865 870 875
 Glu Lys Leu Gln Leu Gln Glu Gln Leu Gln Ala Lys Thr Glu Leu Cys
 885 890 895
 Ala Glu Ala Glu Glu Leu Arg Ala Arg Leu Thr Ala Lys Lys Gln Glu
 900 905 910
 Leu Glu Glu Ile Cys His Asp Leu Glu Ala Arg Val Glu Glu Glu Glu
 915 920 925
 Glu Arg Cys Gln Tyr Leu Gln Ala Glu Lys Lys Lys Met Gln Gln Asn
 930 935 940
 Ile Gln Glu Leu Glu Glu Gln Leu Glu Glu Glu Ser Ala Arg Gln
 945 950 955 960
 Lys Leu Gln Leu Glu Lys Val Thr Thr Glu Ala Lys Leu Lys Lys Leu
 965 970 975
 Glu Glu Asp Gln Ile Ile Met Glu Asp Gln Asn Cys Lys Leu Ala Lys
 980 985 990
 Glu Lys Lys Leu Leu Glu Asp Arg Val Ala Glu Phe Thr Thr Asp Leu
 995 1000 1005
 Met Glu Glu Glu Glu Lys Ser Lys Ser Leu Ala Lys Leu Lys Asn Lys
 1010 1015 1020
 His Glu Ala Met Ile Thr Asp Leu Glu Glu Arg Leu Arg Arg Glu Glu
 1025 1030 1035 1040
 Lys Gln Arg Gln Glu Leu Glu Lys Thr Arg Arg Lys Leu Glu Gly Asp
 1045 1050 1055
 Ser Thr Asp Leu Ser Asp Gln Ile Ala Glu Leu Gln Ala Gln Ile Ala
 1060 1065 1070
 Glu Leu Lys Met Gln Leu Ala Lys Lys Glu Glu Glu Leu Gln Ala Ala
 1075 1080 1085
 Leu Ala Arg Val Glu Glu Glu Ala Ala Gln Lys Asn Met Ala Leu Lys
 1090 1095 1100
 Lys Ile Arg Glu Leu Glu Thr Gln Ile Ser Glu Leu Gln Glu Asp Leu
 1105 1110 1115 1120
 Glu Ser Glu Arg Ala Cys Arg Asn Lys Ala Glu Lys Gln Lys Arg Asp
 1125 1130 1135
 Leu Gly Glu Glu Leu Glu Ala Leu Lys Thr Glu Leu Glu Asp Thr Leu
 1140 1145 1150
 Asp Ser Thr Ala Ala Gln Gln Glu Leu Arg Ser Lys Arg Glu Gln Glu
 1155 1160 1165
 Val Ser Ile Leu Lys Lys Thr Leu Glu Asp Glu Ala Lys Thr His Glu
 1170 1175 1180

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Ala Gln Ile Gln Glu Met Arg Gln Lys His Ser Gln Ala Val Glu Glu
 1185 1190 1195 1200
 Leu Ala Glu Gln Leu Glu Gln Thr Lys Arg Val Lys Ala Thr Leu Glu
 1205 1210 1215
 Lys Ala Lys Gln Thr Leu Glu Asn Glu Arg Gly Glu Leu Ala Asn Glu
 1220 1225 1230
 Val Lys Ala Leu Leu Gln Gly Lys Gly Asp Ser Glu His Lys Arg Lys
 1235 1240 1245
 Lys Val Glu Ala Gln Leu Gln Glu Leu Gln Val Lys Phe Ser Glu Gly
 1250 1255 1260
 Glu Arg Val Arg Thr Glu Leu Ala Asp Lys Val Ser Lys Leu Gln Val
 1265 1270 1275 1280
 Glu Leu Asp Ser Val Thr Gly Leu Leu Asn Gln Ser Asp Ser Lys Ser
 1285 1290 1295
 Ser Lys Leu Thr Lys Asp Phe Ser Ala Leu Glu Ser Gln Leu Gln Asp
 1300 1305 1310
 Thr Gln Glu Leu Leu Gln Glu Glu Asn Arg Gln Lys Leu Ser Leu Ser
 1315 1320 1325
 Thr Lys Leu Lys Gln Met Glu Asp Glu Lys Asn Ser Phe Arg Glu Gln
 1330 1335 1340
 Leu Glu Glu Glu Glu Glu Glu Ala Lys Arg Asn Leu Glu Lys Gln Ile
 1345 1350 1355 1360
 Ala Thr Leu His Ala Gln Val Thr Asp Met Lys Lys Lys Met Glu Asp
 1365 1370 1375
 Gly Val Gly Cys Leu Glu Thr Ala Glu Glu Ala Lys Arg Arg Leu Gln
 1380 1385 1390
 Lys Asp Leu Glu Gly Leu Ser Gln Arg Leu Glu Glu Lys Val Ala Ala
 1395 1400 1405
 Tyr Asp Lys Leu Glu Lys Thr Lys Thr Arg Leu Gln Gln Glu Leu Asp
 1410 1415 1420
 Asp Leu Leu Val Asp Leu Asp His Gln Arg Gln Ser Val Ser Asn Leu
 1425 1430 1435 1440
 Glu Lys Lys Gln Lys Lys Phe Asp Gln Leu Leu Ala Glu Glu Lys Thr
 1445 1450 1455
 Ile Ser Ala Lys Tyr Ala Glu Glu Arg Asp Arg Ala Glu Ala Glu Ala
 1460 1465 1470
 Arg Glu Lys Glu Thr Lys Ala Leu Ser Leu Ala Arg Ala Leu Glu Glu
 1475 1480 1485
 Ala Met Glu Gln Lys Ala Glu Leu Glu Arg Leu Asn Lys Gln Phe Arg
 1490 1495 1500
 Thr Glu Met Glu Asp Leu Met Ser Ser Lys Asp Asp Val Gly Lys Ser
 1505 1510 1515 1520

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Val His Glu Leu Glu Lys Ser Asn Arg Ala Leu Glu Gln Gln Val Glu
 1525 1530 1535
 Glu Met Lys Thr Gln Leu Glu Glu Leu Glu Asp Glu Leu Gln Ala Thr
 1540 1545 1550
 Glu Asp Ala Lys Leu Arg Leu Glu Val Asn Leu Gln Ala Met Lys Ala
 1555 1560 1565
 Gln Phe Glu Arg Asp Leu Gln Gly Arg Asp Glu Gln Ser Glu Glu Lys
 1570 1575 1580
 Lys Lys Gln Leu Val Arg Gln Val Arg Glu Met Glu Ala Glu Leu Glu
 1585 1590 1595 1600
 Asp Glu Arg Lys Gln Arg Ser Ile Ala Met Ala Ala Arg Lys Lys Leu
 1605 1610 1615
 Glu Met Asp Leu Lys Asp Leu Glu Ala His Ile Asp Thr Ala Asn Lys
 1620 1625 1630
 Asn Arg Glu Glu Ala Ile Lys Gln Leu Arg Lys Leu Gln Ala Gln Met
 1635 1640 1645
 Lys Asp Cys Met Arg Asp Val Asp Asp Thr Arg Ala Ser Arg Glu Glu
 1650 1655 1660
 Ile Leu Ala Gln Ala Lys Glu Asn Glu Lys Lys Leu Lys Ser Met Glu
 1665 1670 1675 1680
 Ala Glu Met Ile Gln Leu Gln Glu Glu Leu Ala Ala Ala Glu Arg Ala
 1685 1690 1695
 Lys Arg Gln Ala Gln Gln Glu Arg Asp Glu Leu Ala Asp Glu Ile Ala
 1700 1705 1710
 Asn Ser Ser Gly Lys Gly Ala Leu Ala Leu Glu Glu Lys Arg Arg Leu
 1715 1720 1725
 Glu Ala Leu Ile Ala Leu Leu Glu Glu Glu Leu Glu Glu Gln Gly
 1730 1735 1740
 Asn Thr Glu Leu Ile Asn Asp Arg Leu Lys Lys Ala Asn Leu Gln Ile
 1745 1750 1755 1760
 Asp Gln Ile Asn Thr Asp Leu Asn Leu Glu Arg Ser His Ala Gln Lys
 1765 1770 1775
 Asn Glu Asn Ala Arg Gln Gln Leu Glu Arg Gln Asn Lys Glu Leu Lys
 1780 1785 1790
 Ala Lys Leu Gln Glu Met Glu Ser Ala Val Lys Ser Lys Tyr Lys Ala
 1795 1800 1805
 Ser Ile Ala Ala Leu Glu Ala Lys Ile Ala Gln Leu Glu Glu Gln Leu
 1810 1815 1820
 Asp Asn Glu Thr Lys Glu Arg Gln Ala Ala Ser Lys Gln Val Arg Arg
 1825 1830 1835 1840
 Ala Glu Lys Lys Leu Lys Asp Val Leu Leu Gln Val Glu Asp Glu Arg
 1845 1850 1855

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Arg Asn Ala Glu Gln Phe Lys Asp Gln Ala Asp Lys Ala Ser Thr Arg
1860 1865 1870
Leu Lys Gln Leu Lys Arg Gln Leu Glu Glu Ala Glu Glu Glu Ala Gln
1875 1880 1885
Arg Ala Asn Ala Ser Arg Arg Lys Leu Gln Arg Glu Leu Glu Asp Ala
1890 1895 1900
Thr Glu Thr Ala Asp Ala Met Asn Arg Glu Val Ser Ser Leu Lys Asn
1905 1910 1915 1920
Lys Leu Arg Arg Gly Asp Met Pro Phe Val Val Thr Arg Arg Ile Val
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35 40 45
Glu Val Gly Asp Glu Ala Ile Val Glu Leu Ala Glu Asn Gly Lys Lys
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Val Lys Val Asn Lys Asp Asp Ile Gln Lys Met Asn Pro Pro Lys Phe
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Ser Lys Val Glu Asp Met Ala Glu Leu Thr Cys Leu Asn Glu Ala Ser
85 90 95
Val Leu His Asn Leu Lys Glu Arg Tyr Tyr Ser Gly Leu Ile Tyr Thr
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Tyr Ser Gly Leu Phe Cys Val Val Ile Asn Pro Tyr Lys Asn Leu Pro
115 120 125
Ile Tyr Ser Glu Glu Ile Val Glu Met Tyr Lys Gly Lys Lys Arg His
130 135 140
Glu Met Pro Pro His Ile Tyr Ala Ile Thr Asp Thr Ala Tyr Arg Ser
145 150 155 160
Met Met Gln Asp Arg Glu Asp Gln Ser Ile Leu Cys Thr Gly Glu Ser
165 170 175
Gly Ala Gly Lys Thr Glu Asn Thr Lys Lys Val Ile Gln Tyr Leu Ala
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180					185					190					
His	Val	Ala	Ser	Ser	His	Lys	Ser	Lys	Lys	Asp	Gln	Gly	Glu	Leu	Glu
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Arg	Gln	Leu	Leu	Gln	Ala	Asn	Pro	Ile	Leu	Glu	Ala	Phe	Gly	Asn	Ala
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Lys	Thr	Val	Lys	Asn	Asp	Asn	Ser	Ser	Arg	Phe	Gly	Lys	Phe	Ile	Arg
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Ile	Asn	Phe	Asp	Val	Asn	Gly	Tyr	Ile	Val	Gly	Ala	Asn	Ile	Glu	Thr
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			260					265					270		
Thr	Phe	His	Ile	Phe	Tyr	Tyr	Leu	Leu	Ser	Gly	Ala	Gly	Glu	His	Leu
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Phe	Thr	Arg	Gly	Ile	Leu	Thr	Pro	Arg	Ile	Lys	Val	Gly	Arg	Asp	Tyr
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Val	Gln	Lys	Ala	Gln	Thr	Lys	Glu	Gln	Ala	Asp	Phe	Ala	Ile	Glu	Ala
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			420					425					430		
Ile	Asn	Lys	Ala	Leu	Asp	Lys	Thr	Lys	Arg	Gln	Gly	Ala	Ser	Phe	Ile
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Gly	Ile	Leu	Asp	Ile	Ala	Gly	Phe	Glu	Ile	Phe	Glu	Leu	Asn	Ser	Phe
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Glu	Gln	Leu	Cys	Ile	Asn	Tyr	Thr	Asn	Glu	Lys	Leu	Gln	Gln	Leu	Phe
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Asn	His	Thr	Met	Phe	Ile	Leu	Glu	Gln	Glu	Glu	Tyr	Gln	Asn	Glu	Gly
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Asp	Leu	Ile	Glu	Lys	Pro	Ala	Gly	Pro	Pro	Gly	Ile	Leu	Ala	Leu	Leu

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Lys	Val	Val	Gln	Glu	Gln	Gly	Thr	His	Pro	Lys	Phe	Gln	Lys	Pro	Lys
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Gln	Leu	Lys	Asp	Lys	Ala	Asp	Phe	Cys	Ile	Ile	His	Tyr	Ala	Gly	Lys
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Asn	Asp	Asn	Ile	Ala	Thr	Leu	Leu	His	Gln	Ser	Ser	Asp	Lys	Phe	Val
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Ser	Glu	Leu	Trp	Lys	Asp	Val	Asp	Arg	Ile	Val	Gly	Leu	Asp	Gln	Val
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Gly	Met	Phe	Arg	Thr	Val	Gly	Gln	Leu	Tyr	Lys	Glu	Gln	Leu	Ala	Lys
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Leu	Met	Ala	Thr	Leu	Arg	Asn	Thr	Asn	Pro	Asn	Phe	Val	Arg	Cys	Ile
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Ile	Pro	Asn	His	Glu	Lys	Lys	Ala	Gly	Lys	Leu	Asp	Pro	His	Leu	Val
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Leu	Asp	Gln	Leu	Arg	Cys	Asn	Gly	Val	Leu	Glu	Gly	Ile	Arg	Ile	Cys
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Tyr	Glu	Ile	Leu	Thr	Pro	Asn	Ala	Ile	Pro	Lys	Gly	Phe	Met	Asp	Gly
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Lys	Gln	Ala	Cys	Val	Leu	Met	Ile	Lys	Ala	Leu	Glu	Leu	Asp	Ser	Asn
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Ala	His	Leu	Glu	Glu	Glu	Arg	Asp	Leu	Lys	Ile	Thr	Asp	Val	Ile	Ile
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Lys	Val	Lys	Pro	Leu	Leu	Gln	Val	Ser	Arg	Gln	Glu	Glu	Glu	Met	Met
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 Glu Lys Met Gln Leu Gln Glu Gln Leu Gln Ala Glu Ala Glu Leu Cys
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 Ile Gln Glu Leu Glu Glu Gln Leu Glu Glu Glu Glu Ser Ala Arg Gln
 945 950 955 960
 Lys Leu Gln Leu Glu Lys Val Thr Thr Glu Ala Lys Leu Lys Lys Leu
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 995 1000 1005
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 His Glu Ala Met Ile Thr Asp Leu Glu Glu Arg Leu Arg Arg Glu Glu
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 Val Lys Val Leu Leu Gln Gly Lys Gly Asp Ala Glu His Lys Arg Lys
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 Lys Val Asp Ala Gln Leu Gln Glu Leu Gln Val Lys Phe Thr Glu Gly
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 1265 1270 1275 1280
 Glu Leu Asp Asn Val Thr Gly Leu Leu Asn Gln Ser Asp Ser Lys Ser
 1285 1290 1295
 Ile Lys Leu Ala Lys Asp Phe Ser Ala Leu Glu Ser Gln Leu Gln Asp
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 Thr Gln Glu Leu Leu Gln Glu Glu Thr Arg Leu Lys Leu Ser Phe Ser
 1315 1320 1325
 Thr Lys Leu Lys Gln Thr Glu Asp Glu Lys Asn Ala Leu Lys Glu Gln
 1330 1335 1340
 Leu Glu Glu Glu Glu Glu Ala Lys Arg Asn Leu Glu Lys Gln Ile Ser
 1345 1350 1355 1360
 Val Leu Gln Gln Gln Ala Val Glu Ala Arg Lys Lys Met Asp Asp Gly
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 Leu Gly Cys Leu Glu Ile Ala Glu Glu Ala Lys Lys Lys Leu Gln Lys
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 1395 1400 1405
 Asp Lys Leu Glu Lys Thr Lys Thr Arg Leu Gln Gln Glu Leu Asp Asp
 1410 1415 1420
 Ile Ala Val Asp Leu Asp His Gln Arg Gln Thr Val Ser Asn Leu Glu
 1425 1430 1435 1440
 Lys Lys Gln Lys Lys Phe Asp Gln Leu Leu Ala Glu Glu Lys Asn Ile
 1445 1450 1455
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 Glu Lys Glu Thr Lys Ala Leu Ser Leu Ala Arg Ala Leu Glu Glu Ala
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 1490 1495 1500
 Glu Met Glu Asp Leu Met Ser Ser Lys Asp Asp Val Gly Lys Ser Val
 1505 1510 1515 1520
 His Glu Leu Glu Lys Ala Lys Arg Ala Leu Glu Gln Gln Val Glu Glu
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1525		1535
Met Lys Thr Gln Leu Glu Glu Leu Glu Asp Glu Leu Gln Ala Thr Glu		
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Asp Ala Lys Leu Arg Leu Glu Val Asn Gln Gln Ala Met Lys Ala Gln		
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Phe Asp Arg Asp Leu Leu Gly Arg Asp Glu Gln Asn Glu Glu Lys Arg		
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Lys Gln Leu Ile Arg Gln Val Arg Glu Met Glu Val Glu Leu Glu Asp		
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Glu Arg Lys Gln Arg Ser Ile Ala Val Ala Ala Arg Lys Lys Leu Glu		
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Leu Asp Leu Lys Asp Leu Glu Ser His Ile Asp Thr Ala Asn Lys Asn		
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Arg Asp Glu Ala Ile Lys His Val Arg Lys Leu Gln Ala Gln Met Lys		
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Asp Tyr Met Arg Glu Leu Glu Asp Thr Arg Thr Ser Arg Glu Glu Ile		
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Leu Ala Gln Ala Lys Glu Asn Glu Lys Lys Leu Lys Ser Met Glu Ala		
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Ser Ser Gly Lys Gly Ala Leu Ala Met Glu Glu Lys Arg Arg Leu Glu		
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Ala Arg Ile Ala Gln Leu Glu Glu Glu Leu Glu Glu Glu Gln Gly Asn		
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Thr Glu Ile Ile Asn Asp Arg Leu Lys Lys Ala Asn Leu Gln Ile Asp		
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Gln Met Asn Ala Asp Leu Asn Ala Glu Arg Ser Asn Ala Gln Lys Asn		
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Glu Asn Ala Arg Gln Gln Met Glu Arg Gln Asn Lys Glu Leu Lys Leu		
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Lys Leu Gln Glu Met Glu Ser Ala Val Lys Ser Lys Tyr Lys Ala Thr		
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Ile Thr Ala Leu Glu Ala Lys Ile Val Gln Leu Glu Glu Gln Leu Asp		
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Glu Lys Lys Leu Lys Asp Ile Leu Leu Gln Val Asp Asp Glu Arg Arg		
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Asn Ala Glu Gln Phe Lys Asp Gln Ala Asp Lys Ala Asn Met Arg Leu		

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 Val Lys Val Asn Lys Asp Asp Ile Gln Lys Met Asn Pro Pro Lys Phe
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 Ser Lys Val Glu Asp Met Ala Glu Leu Thr Cys Leu Asn Glu Ala Ser
 85 90 95
 Val Leu His Asn Leu Lys Glu Arg Tyr Tyr Ser Gly Leu Ile Tyr Thr
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 Tyr Ser Gly Leu Phe Cys Val Val Ile Asn Pro Tyr Lys Asn Leu Pro
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 Met Met Gln Asp Arg Glu Asp Gln Ser Ile Leu Cys Thr Gly Glu Ser
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 Gly Ala Gly Lys Thr Glu Asn Thr Lys Lys Val Ile Gln Tyr Leu Ala
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Arg	Gln	Leu	Leu	Gln	Ala	Asn	Pro	Ile	Leu	Glu	Ala	Phe	Gly	Asn	Ala		
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Ile	Asn	Phe	Asp	Val	Asn	Gly	Tyr	Ile	Val	Gly	Ala	Asn	Ile	Glu	Thr		
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Tyr	Leu	Leu	Glu	Lys	Ser	Arg	Ala	Ile	Arg	Gln	Ala	Lys	Glu	Glu	Arg		
			260					265					270				
Thr	Phe	His	Ile	Phe	Tyr	Tyr	Leu	Leu	Ser	Gly	Ala	Gly	Glu	His	Leu		
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Lys	Thr	Asp	Leu	Leu	Leu	Glu	Pro	Tyr	Gly	Lys	Tyr	Arg	Phe	Leu	Ser		
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Thr	Ala	Ala	Gln	Lys	Val	Ser	His	Leu	Leu	Gly	Ile	Asn	Val	Thr	Asp		
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Glu	Trp	Asn	Phe	Ile	Asp	Phe	Gly	Leu	Asp	Leu	Gln	Pro	Cys	Ile	Asp		
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Leu	Ile	Glu	Arg	Pro	Ala	Asn	Pro	Pro	Gly	Val	Leu	Ala	Leu	Leu	Asp		
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Glu	Leu	Trp	Lys	Asp	Val	Asp	Arg	Ile	Val	Gly	Leu	Asp	Gln	Val	Thr
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Gly	Ile	Thr	Glu	Thr	Ala	Phe	Gly	Ser	Ala	Tyr	Lys	Thr	Lys	Lys	Gly
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Asp	Gln	Leu	Arg	Cys	Asn	Gly	Val	Leu	Glu	Gly	Ile	Arg	Ile	Cys	Arg
	690					695					700				
Gln	Gly	Phe	Pro	Asn	Arg	Ile	Val	Phe	Gln	Glu	Phe	Arg	Gln	Arg	Tyr
705					710					715					720
Glu	Ile	Leu	Thr	Pro	Asn	Ala	Ile	Pro	Lys	Gly	Phe	Met	Asp	Gly	Lys
				725					730					735	
Gln	Ala	Cys	Glu	Arg	Met	Ile	Arg	Ala	Leu	Glu	Leu	Asp	Pro	Asn	Leu
			740					745					750		
Tyr	Arg	Ile	Gly	Gln	Ser	Lys	Ile	Phe	Phe	Arg	Ala	Gly	Val	Leu	Ala
		755					760					765			
His	Leu	Glu	Glu	Glu	Arg	Asp	Leu	Lys	Ile	Thr	Asp	Ile	Ile	Ile	Phe
	770					775					780				
Phe	Gln	Ala	Val	Cys	Arg	Gly	Tyr	Leu	Ala	Arg	Lys	Ala	Phe	Ala	Lys
785					790					795					800
Lys	Gln	Gln	Gln	Leu	Ser	Ala	Leu	Lys	Ile	Leu	Gln	Arg	Asn	Cys	Ala
				805					810					815	
Ala	Tyr	Leu	Lys	Leu	Arg	His	Trp	Gln	Trp	Trp	Arg	Val	Phe	Thr	Lys
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Val	Lys	Pro	Leu	Leu	Gln	Val	Thr	Arg	Gln	Glu	Glu	Glu	Leu	Gln	Ala
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Lys Asn Ile Leu Ala Glu Gln Leu Gln Ala Glu Thr Glu Leu Phe Ala
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Glu Ala Glu Glu Met Arg Ala Arg Leu Ala Ala Lys Lys Gln Glu Leu
900 905 910

Glu Glu Ile Leu His Asp Leu Glu Ser Arg Val Glu Glu Glu Glu Glu
915 920 925

Arg Asn Gln Ile Leu Gln Asn Glu Lys Lys Lys Glu Gln Gly His Lys
930 935 940

Asn Asp Leu Glu Glu Gln Leu Asp Glu Met Glu Ser Ala Arg Gln Lys
945 950 955 960

Leu Gln Leu Glu Lys Val Thr Thr Glu Ala Lys Leu Lys Lys Leu Glu
965 970 975

Glu Glu Gln Ile Ile Leu Glu Asp Gln Asn Cys Lys Leu Ala Lys Glu
980 985 990

Lys Lys Leu Leu Glu Asp Arg Ile Ala Glu Phe Thr Thr Asn Leu Thr
995 1000 1005

Glu Glu Glu Glu Lys Ser Lys Ser Leu Ala Lys Leu Lys Asn Lys His
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Glu Ala Met Ile Thr Asp Leu Glu Glu Arg Leu Arg Arg Glu Glu Lys
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Gln Arg Gln Glu Leu Glu Lys Thr Arg Arg Lys Leu Glu Gly Asp Ser
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Thr Asp Leu Ser Asp Gln Ile Ala Glu Leu Gln Ala Gln Ile Ala Glu
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Leu Lys Met Gln Leu Ala Lys Lys Glu Glu Glu Leu Gln Ala Ala Leu
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Ala Arg Val Glu Glu Glu Ala Ala Gln Lys Asn Met Ala Leu Lys Lys
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Ser Glu Arg Ala Ser Arg Asn Lys Ala Glu Lys Gln Lys Arg Asp Leu
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Gly Glu Glu Leu Glu Ala Leu Lys Thr Glu Leu Glu Asp Leu Thr Asp
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Ser Thr Ala Ala Gln Gln Glu Leu Arg Ser Lys Arg Glu Gln Glu Val
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Asn Ile Leu Lys Lys Thr Leu Glu Glu Glu Ala Lys Thr His Glu Ala
1170 1175 1180

Gln Ile Gln Glu Met Arg Gln Lys His Ser Gln Ala Val Glu Glu Leu
1185 1190 1195 1200

Ala Glu Gln Leu Glu Gln Thr Lys Arg Lys Val Ala Asn Leu Glu Lys
 1205 1210 1215
 Ala Lys Gln Thr Leu Glu Asn Glu Arg Gly Glu Leu Ala Asn Glu Val
 1220 1225 1230
 Lys Val Leu Leu Gln Gly Gly Arg Asp Ser Glu His Lys Arg Lys Lys
 1235 1240 1245
 Val Glu Ala Gln Leu Gln Glu Leu Gln Val Lys Phe Asn Glu Gly Glu
 1250 1255 1260
 Arg Arg Val Thr Glu Leu Ala Asp Lys Val Thr Lys Leu Gln Val Glu
 1265 1270 1275 1280
 Leu Asp Asn Val Thr Gly Leu Leu Ser Gln Ser Asp Ser Lys Ser Ser
 1285 1290 1295
 Lys Leu Thr Lys Asp Phe Ser Ala Leu Glu Ser Gln Leu Gln Asp Thr
 1300 1305 1310
 Gln Glu Leu Leu Gln Glu Glu Asn Arg Gln Lys Leu Ser Leu Ser Thr
 1315 1320 1325
 Lys Leu Lys Gln Val Glu Asp Glu Lys Asn Ser Phe Arg Glu Gln Leu
 1330 1335 1340
 Glu Glu Glu Glu Glu Glu Ala Lys His Asn Leu Glu Lys Gln Ile Ala
 1345 1350 1355 1360
 Thr Leu His Ala Gln Val Ala Asp Met Lys Lys Lys Met Glu Asp Ser
 1365 1370 1375
 Val Gly Cys Leu Glu Thr Ala Glu Glu Val Lys Arg Lys Leu Gln Lys
 1380 1385 1390
 Asp Leu Glu Gly Leu Ser Gln Arg His Glu Glu Lys Val Ala Ala Tyr
 1395 1400 1405
 Asp Lys Leu Glu Lys Thr Lys Thr Arg Leu Gln Gln Glu Leu Asp Asp
 1410 1415 1420
 Leu Leu Val Asp Leu Asp His Gln Arg Gln Ser Ala Cys Asn Leu Glu
 1425 1430 1435 1440
 Lys Lys Gln Lys Lys Phe Asp Gln Leu Leu Ala Glu Glu Ile Thr Lys
 1445 1450 1455
 Ser Ala Lys Tyr Ala Glu Glu Arg Ala Arg Asp Ala Glu Glu Arg Ala
 1460 1465 1470
 Glu Lys Ala Thr Lys Glu Leu Ser Leu Ala Arg Ala Glu Leu Glu Ala
 1475 1480 1485
 Met Glu Gln Lys Ala Glu Phe Leu Arg Lys Asn Leu Gln Glu Met Thr
 1490 1495 1500
 Glu Arg Leu Asp Glu Met Ser Ser Lys Val Asp Asp Ala Lys Ser Val
 1505 1510 1515 1520
 Leu Glu His Glu Lys Ser Lys Leu Gly Arg Glu Gln Gln Val Met Glu
 1525 1530 1535

Glu Lys Thr Gln Leu Leu Glu Glu Glu Asp Glu Leu Ala Gln Thr Glu
 1540 1545 1550
 Asp Ala Lys Leu Arg Leu Glu Val Asn Leu Gln Ala Met Lys Ala Gln
 1555 1560 1565
 Phe Glu Arg Asp Leu Gln Gly Arg Gln Asp Asp Ser Glu Glu Lys Gln
 1570 1575 1580
 Lys Lys Leu Val Arg Gln Val Arg Glu Met Glu Ala Glu Leu Glu Asp
 1585 1590 1595 1600
 Gln Arg Lys Glu Met Ser Arg Ala Arg Ala Ala Val Lys Lys Leu Glu
 1605 1610 1615
 Met Asp Leu Lys Asp Leu Glu Ala His Ile Asp Ser Ala Asn Lys Asn
 1620 1625 1630
 Arg Asp Glu Ala Lys Ile Gln Leu Arg Asn Leu Gln Ala Gln Met Lys
 1635 1640 1645
 Asp Cys Met Arg Glu Leu Asp Asp Thr Arg Ala Ser Arg Glu Glu Ile
 1650 1655 1660
 Ala Leu Gln Ala Lys Glu Asn Glu Lys Lys Leu Lys Ser Met Glu Ala
 1665 1670 1675 1680
 Glu Met Ile Gln Leu Gln Glu Glu Leu Ala Ala Ala Glu Arg Ala Lys
 1685 1690 1695
 Arg Gln Ala Gln Gln Glu Arg Asp Glu Leu Ala Asp Glu Ile Ser Asn
 1700 1705 1710
 Ala Ser Gly Lys Ala Gly Leu Ala Lys Glu Glu Leu Arg Arg Leu Glu
 1715 1720 1725
 Ala Arg Ile Ala Gln Leu Glu Glu Glu Leu Glu Glu Gln Gly Asn
 1730 1735 1740
 Thr Glu Leu Ile Asn Asp Arg Leu Lys Lys Ala Asn Leu Gln Ile Asp
 1745 1750 1755 1760
 Gln Ile Asn Ala Asp Leu Asn Leu Glu Arg Gly His Ala Gln Lys Asn
 1765 1770 1775
 Glu Asn Ala Arg Gln Gln Leu Glu Arg Gln Asn Lys Glu Leu Lys Val
 1780 1785 1790
 Lys Leu Gln Glu Met Glu Gly Thr Val Lys Ser Lys Tyr Lys Ala Ser
 1795 1800 1805
 Ile Thr Ala Leu Glu Ala Lys Ile Ala Gln Leu Glu Glu Gln Leu Asp
 1810 1815 1820
 Asn Glu Thr Lys Glu Arg Gln Ala Ala Cys Lys Gln Val Arg Arg Thr
 1825 1830 1835 1840
 Glu Lys Lys Leu Lys Asp Val Leu Leu Gln Val Asp Asp Glu Arg Arg
 1845 1850 1855
 Asn Ala Glu Gln Tyr Lys Asp Gln Ala Asp Lys Ala Ser Thr Arg Leu
 1860 1865 1870

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Lys Gln Leu Lys Arg Gln Leu Glu Glu Ala Glu Glu Glu Ala Gln Arg
1875 1880 1885

Ala Asn Ala Ser Arg Arg Lys Leu Gln Arg Glu Leu Glu Asp Ala Thr
1890 1895 1900

Glu Thr Ala Asp Ala Met Asn Arg Glu Val Ser Ser Leu Lys Asn Lys
1905 1910 1915 1920

Leu Arg Arg Gly Asp Leu Pro Phe Val Val Thr Arg Arg Leu Val Arg
1925 1930 1935

Lys Gly Thr Leu Glu Leu Ser Asp Asp Asp Asp Glu Ser Lys Ala Ser
1940 1945 1950

Leu Ile Asn Glu Thr Gln Pro Pro Gln Cys Leu Asp Gln Gln Leu Asp
1955 1960 1965

Gln Leu Phe His Trp Pro Val Asn Ala Gly Cys Val Cys Gly Trp Gly
1970 1975 1980

Val Glu Gln Thr Gln Gly Glu Glu Ala Val His Lys Cys Arg Thr
1985 1990 1995

<210> 108

<211> 734

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Myosin Head
(motor domain) sequence

<400> 108

Val Glu Asp Met Val Glu Leu Thr Tyr Leu Asn Glu Pro Ser Val Leu
1 5 10 15

His Asn Leu Lys Lys Arg Tyr Lys Ser Asp Leu Ile Tyr Thr Tyr Ser
20 25 30

Gly Leu Val Leu Val Ser Val Asn Pro Tyr Lys Arg Leu Pro Gln Ile
35 40 45

Tyr Thr Glu Glu Ile Ile Ala Lys Tyr Arg Gly Lys Arg Arg Tyr Glu
50 55 60

Leu Pro Pro His Ile Phe Ala Ile Ala Asp Glu Ala Tyr Arg Ser Met
65 70 75 80

Leu Ser Asp Lys Glu Asn Gln Ser Ile Leu Ile Ser Gly Glu Ser Gly
85 90 95

Ala Gly Lys Thr Glu Asn Thr Lys Lys Val Met Gln Tyr Leu Ala Ala
100 105 110

Val Ser Gly Gly Asn Ser Gly Asn Gly Glu Glu Val Pro Ser Val Lys
115 120 125

Val Gly Arg Val Glu Asp Gln Ile Leu Gln Ser Asn Pro Ile Leu Glu
130 135 140

Ala Phe Gly Asn Ala Lys Thr Thr Arg Asn Asn Asn Ser Ser Arg Phe
Page 260

145		150		155		160									
Gly	Lys	Tyr	Ile	Glu	Ile	Gln	Phe	Asp	Lys	Thr	Gly	Lys	Ile	Val	Gly
				165					170					175	
Ala	Lys	Ile	Glu	Asn	Tyr	Leu	Leu	Glu	Lys	Ser	Arg	Val	Val	Tyr	Gln
			180					185					190		
Thr	Glu	Gly	Glu	Arg	Asn	Phe	His	Ile	Phe	Tyr	Gln	Leu	Leu	Ala	Gly
		195					200					205			
Ala	Ser	Gln	Gln	Asn	Leu	Lys	Lys	Glu	Leu	Lys	Leu	Thr	Asn	Asp	Pro
	210					215					220				
Glu	Asp	Tyr	His	Tyr	Leu	Asn	Gln	Gly	Gly	Glu	Val	Lys	Pro	Cys	Tyr
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Thr	Val	Asp	Gly	Ile	Asp	Asp	Ser	Glu	Gly	Asn	Val	Glu	Glu	Phe	Lys
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Glu	Thr	Arg	Lys	Ala	Met	Asp	Ile	Leu	Gly	Phe	Thr	Asp	Glu	Glu	Gln
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Arg	Ser	Ile	Phe	Arg	Ile	Val	Ala	Ala	Ile	Leu	His	Leu	Gly	Asn	Ile
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Lys	Phe	Lys	Gln	Arg	Arg	Lys	Glu	Glu	Ala	Ala	Ile	Pro	Asp	Asp	Asn
	290					295					300				
Asn	Ala	Asp	Thr	Lys	Ala	Leu	Glu	Lys	Ala	Ala	Glu	Leu	Leu	Gly	Val
305					310				315					320	
Asp	Ala	Thr	Glu	Leu	Glu	Lys	Ala	Leu	Leu	Ser	Arg	Arg	Ile	Lys	Thr
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Gly	Thr	Glu	Gly	Arg	Lys	Ser	Thr	Val	Thr	Lys	Pro	Gln	Asn	Val	Glu
			340					345					350		
Gln	Ala	Ser	Tyr	Ala	Arg	Asp	Ala	Leu	Ala	Lys	Ala	Leu	Tyr	Ser	Arg
		355					360					365			
Leu	Phe	Asp	Trp	Ile	Val	Asn	Arg	Ile	Asn	Lys	Thr	Leu	Asp	Phe	Lys
	370					375					380				
Ala	Lys	Glu	Gly	Gln	Asp	Ala	Ser	Phe	Ile	Gly	Val	Leu	Asp	Ile	Tyr
385					390					395				400	
Gly	Phe	Glu	Ile	Phe	Glu	Lys	Asn	Ser	Phe	Glu	Gln	Leu	Cys	Ile	Asn
				405					410					415	
Tyr	Val	Asn	Glu	Lys	Leu	Gln	Gln	Phe	Phe	Asn	His	His	Met	Phe	Lys
			420					425					430		
Leu	Glu	Gln	Glu	Glu	Tyr	Lys	Arg	Glu	Gly	Ile	Glu	Trp	Thr	Phe	Ile
		435					440					445			
Asp	Phe	Gly	Asp	Asn	Leu	Gln	Pro	Cys	Ile	Asp	Leu	Ile	Glu	Lys	Lys
	450					455					460				
Ser	Pro	Pro	Gly	Ile	Leu	Ser	Leu	Leu	Asp	Glu	Glu	Cys	Leu	Phe	Pro
465					470					475				480	
Lys	Ala	Gln	Ser	Gly	Thr	Asp	Gln	Thr	Phe	Leu	Asp	Lys	Leu	Tyr	Ser

485 490 495
 Thr Phe Ser Lys His Pro Ala His Phe Glu Lys Phe Ser Pro Arg Phe
 500 505 510
 Arg Gln Lys Lys Ser Gly Ala His Phe Ile Ile Lys His Tyr Ala Gly
 515 520 525
 Asp Val Glu Tyr Asn Val Glu Gly Phe Leu Glu Lys Asn Lys Asp Pro
 530 535 540
 Leu Phe Asp Asp Leu Ile Ser Leu Leu Lys Ser Ser Ser Asn Pro Leu
 545 550 555 560
 Leu Ala Glu Leu Phe Pro Asp Glu Glu Thr Leu Ala Gly Pro Phe Glu
 565 570 575
 Ala Asp Pro Ser Ser Leu Ser Lys Lys Arg Lys Ser Gly Ser Lys Asn
 580 585 590
 Lys Ser Thr Gly Lys Lys Thr Lys Lys Ser Asn Phe Ile Thr Val Gly
 595 600 605
 Ala Gln Phe Lys Glu Ser Leu Asn Glu Leu Met Lys Thr Leu Ser Ser
 610 615 620
 Thr Asn Leu Pro His Phe Val Arg Cys Ile Lys Pro Asn Glu Lys Lys
 625 630 635 640
 Lys Ala Gly Val Phe Asp Ala Ser Leu Val Leu His Gln Leu Arg Cys
 645 650 655
 Leu Gly Val Leu Glu Gly Ile Arg Ile Arg Arg Ala Gly Phe Pro Asn
 660 665 670
 Arg Ile Thr Phe Asp Glu Phe Leu Gln Arg Tyr Arg Ile Leu Ala Pro
 675 680 685
 Lys Thr Trp Pro Lys Trp Ser Gly Asp Ala Lys Lys Gly Glu Lys Asn
 690 695 700
 Glu Ile Val Ala Cys Glu Lys Leu Leu Gln Ser Leu Asn Leu Asp Lys
 705 710 715 720
 Gly Glu Glu Tyr Arg Phe Gly Lys Thr Lys Ile Phe Phe Arg
 725 730

<210> 109
 <211> 175
 <212> PRT
 <213> Homo sapiens

<400> 109
 Met Leu Pro Pro Met Ala Leu Pro Ser Val Ser Trp Met Leu Leu Ser
 1 5 10 15
 Cys Leu Met Leu Leu Ser Gln Val Gln Gly Glu Glu Pro Gln Arg Glu
 20 25 30
 Leu Pro Ser Ala Arg Ile Arg Cys Pro Lys Gly Ser Lys Ala Tyr Gly
 35 40 45

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Ser His Cys Tyr Ala Leu Phe Leu Ser Pro Lys Ser Trp Thr Asp Ala
50 55 60

Asp Leu Ala Cys Gln Lys Arg Pro Ser Gly Asn Leu Val Ser Val Leu
65 70 75 80

Ser Gly Ala Glu Gly Ser Phe Val Ser Ser Leu Val Lys Ser Ile Gly
85 90 95

Asn Ser Tyr Ser Tyr Val Trp Ile Gly Leu His Asp Pro Thr Gln Gly
100 105 110

Thr Glu Pro Asn Gly Glu Gly Trp Glu Trp Ser Ser Ser Asp Val Met
115 120 125

Asn Tyr Phe Ala Trp Glu Arg Asn Pro Ser Thr Ile Ser Ser Pro Gly
130 135 140

His Cys Ala Ser Leu Ser Arg Ser Thr Ala Phe Leu Arg Trp Lys Asp
145 150 155 160

Tyr Asn Cys Asn Val Arg Leu Pro Tyr Val Cys Lys Phe Thr Asp
165 170 175

<210> 110
<211> 175
<212> PRT
<213> Bos taurus

<400> 110
Met Leu Pro Ser Leu Gly Leu Pro Arg Leu Ser Trp Met Leu Leu Ser
1 5 10 15

Cys Leu Met Leu Leu Ser Gln Ile Gln Gly Glu Asn Ser Gln Lys Glu
20 25 30

Leu Pro Ser Ala Arg Ile Ser Cys Pro Ser Gly Ser Met Ala Tyr Arg
35 40 45

Ser His Cys Tyr Ala Leu Phe Lys Thr Pro Lys Thr Trp Met Asp Ala
50 55 60

Asp Ile Ala Cys Gln Lys Arg Pro Ser Gly His Leu Val Ser Val Leu
65 70 75 80

Ser Gly Ala Glu Glu Ser Phe Val Ala Ser Leu Val Arg Asn Asn Leu
85 90 95

Asn Thr Gln Ser Asp Ile Trp Ile Gly Leu His Asp Pro Thr Glu Gly
100 105 110

Ser Glu Ala Asn Ala Gly Gly Trp Glu Trp Ile Ser Asn Asp Val Leu
115 120 125

Asn Tyr Val Ala Trp Glu Thr Asp Pro Ala Ala Ile Ser Ser Pro Gly
130 135 140

Tyr Cys Gly Ser Leu Ser Arg Ser Ser Gly Tyr Leu Lys Trp Arg Asp
145 150 155 160

His Asn Cys Asn Leu Asn Leu Pro Tyr Val Cys Lys Phe Thr Asp
165 170 175

<210> 111
 <211> 175
 <212> PRT
 <213> Rattus norvegicus

<400> 111
 Met Leu His Arg Leu Ala Phe Pro Val Met Ser Trp Met Leu Leu Ser
 1 5 10 15
 Cys Leu Met Leu Leu Ser Gln Val Gln Gly Glu Asp Ser Pro Lys Lys
 20 25 30
 Ile Pro Ser Ala Arg Ile Ser Cys Pro Lys Gly Ser Gln Ala Tyr Gly
 35 40 45
 Ser Tyr Cys Tyr Ala Leu Phe Gln Ile Pro Gln Thr Trp Phe Asp Ala
 50 55 60
 Glu Leu Ala Cys Gln Lys Arg Pro Glu Gly His Leu Val Ser Val Leu
 65 70 75 80
 Asn Val Ala Glu Ala Ser Phe Leu Ala Ser Met Val Lys Asn Thr Gly
 85 90 95
 Asn Ser Tyr Gln Tyr Thr Trp Ile Gly Leu His Asp Pro Thr Leu Gly
 100 105 110
 Gly Glu Pro Asn Gly Gly Gly Trp Glu Trp Ser Asn Asn Asp Ile Met
 115 120 125
 Asn Tyr Val Asn Trp Glu Arg Asn Pro Ser Thr Ala Leu Asp Arg Gly
 130 135 140
 Phe Cys Gly Ser Leu Ser Arg Ser Ser Gly Phe Leu Arg Trp Arg Asp
 145 150 155 160
 Thr Thr Cys Glu Val Lys Leu Pro Tyr Val Cys Lys Phe Thr Gly
 165 170 175

<210> 112
 <211> 175
 <212> PRT
 <213> Mus musculus

<400> 112
 Met Leu Pro Pro Thr Ala Cys Ser Val Met Ser Trp Met Leu Leu Ser
 1 5 10 15
 Cys Leu Met Leu Leu Ser Gln Val Gln Gly Glu Asp Ser Leu Lys Asn
 20 25 30
 Ile Pro Ser Ala Arg Ile Ser Cys Pro Lys Gly Ser Gln Ala Tyr Gly
 35 40 45
 Ser Tyr Cys Tyr Ala Leu Phe Gln Ile Pro Gln Thr Trp Phe Asp Ala
 50 55 60
 Glu Leu Ala Cys Gln Lys Arg Pro Gly Gly His Leu Val Ser Val Leu
 65 70 75 80

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Asn	Ser	Ala	Glu	Ala 85	Ser	Phe	Leu	Ser	Ser 90	Met	Val	Lys	Arg	Thr 95	Gly
Asn	Ser	Tyr	Gln 100	Tyr	Thr	Trp	Ile	Gly 105	Leu	His	Asp	Pro	Thr 110	Leu	Gly
Ala	Glu	Pro 115	Asn	Gly	Gly	Gly	Trp 120	Glu	Trp	Ser	Asn	Asn 125	Asp	Val	Met
Asn	Tyr 130	Phe	Asn	Trp	Glu	Arg 135	Asn	Pro	Ser	Thr	Ala 140	Leu	Asp	Arg	Ala
Phe 145	Cys	Gly	Ser	Leu	Ser 150	Arg	Ala	Ser	Gly	Phe 155	Leu	Lys	Trp	Arg	Asp 160
Met	Thr	Cys	Glu	Val 165	Lys	Leu	Pro	Tyr	Val 170	Cys	Lys	Phe	Thr	Gly 175	

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<210> 113
<211> 174
<212> PRT
<213> Rattus norvegicus
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<400>	113																
Met	Leu	Pro	Arg	Val	Ala	Leu	Thr	Thr	Met	Ser	Trp	Met	Leu	Leu	Ser		
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Ser	Leu	Met	Leu	Leu	Ser	Gln	Val	Gln	Gly	Glu	Asp	Ala	Lys	Glu	Asp		
			20					25					30				
Val	Pro	Thr	Ser	Arg	Ile	Ser	Cys	Pro	Lys	Gly	Ser	Arg	Ala	Tyr	Gly		
		35					40					45					
Ser	Tyr	Cys	Tyr	Ala	Leu	Phe	Ser	Val	Ser	Lys	Ser	Trp	Phe	Asp	Ala		
	50					55					60						
Asp	Leu	Ala	Cys	Gln	Lys	Arg	Pro	Ser	Gly	His	Leu	Val	Ser	Val	Leu		
65					70					75					80		
Ser	Gly	Ser	Glu	Ala	Ser	Phe	Val	Ser	Ser	Leu	Ile	Lys	Ser	Ser	Gly		
				85					90				95				
Asn	Ser	Gly	Gln	Asn	Val	Trp	Ile	Gly	Leu	His	Asp	Pro	Thr	Leu	Gly		
			100					105					110				
Gln	Glu	Pro	Asn	Arg	Gly	Gly	Trp	Glu	Trp	Ser	Asn	Ala	Asp	Val	Met		
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Asn	Tyr	Phe	Asn	Trp	Glu	Thr	Asn	Pro	Ser	Ser	Val	Ser	Gly	Ser	His		
	130					135					140						
Cys	Gly	Thr	Leu	Thr	Arg	Ala	Ser	Gly	Phe	Leu	Arg	Trp	Arg	Glu	Asn		
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Asn	Cys	Ile	Ser	Glu	Leu	Pro	Tyr	Val	Cys	Lys	Phe	Lys	Ala				
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<210> 114
<211> 125
<212> PRT
<213> Artificial Sequence
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<220>

<223> Description of Artificial Sequence: Lectin-C type domain sequence

<400> 114

Glu Ser Lys Thr Trp Ala Glu Ala Glu Leu Ala Cys Gln Lys Glu Gly
 1 5 10 15
 Gly His Ala His Leu Val Ser Ile Gln Ser Ala Glu Glu Gln Ser Phe
 20 25 30
 Val Val Ala Phe Leu Thr Ser Leu Thr Lys Lys Ser Asn Thr Tyr Ala
 35 40 45
 Trp Ile Gly Leu Thr Asp Ile Asn Thr Glu Gly Thr Trp Val Trp Glu
 50 55 60
 Gly Trp Glu Thr Asp Gly Ser Pro Val Asn Tyr Thr Glu Asn Trp Ala
 65 70 75 80
 Pro Gly Glu Pro Asn Asn Arg Gly Asn His Gly Gly Asn Glu Asp Cys
 85 90 95
 Val Glu Ile Tyr Thr Asp Thr Asp Phe Leu Ala Gly Lys Trp Asn Asp
 100 105 110
 Glu Pro Cys Asp Ser Lys Leu Pro Tyr Val Cys Glu Phe
 115 120 125

<210> 115

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer sequence

<400> 115

ctggtttagt gttgccatgg t

21

<210> 116

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer sequence

<400> 116

cagcttcgtt ggcacaggcc tctc

24

<210> 117

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 117
ccagtataag ctgacctttg acaaag 26

<210> 118
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 118
ctggttgtag gttgccatgg t 21

<210> 119
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 119
cagcttcgtt ggcacaggcc tctc 24

<210> 120
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 120
ccagtataag ctgacctttg acaaag 26

<210> 121
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 121
ccaaggtttt agctgtggat ct 22

<210> 122
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 122
 acatccactg cctggaagac cctg 24

<210> 123
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 123
 cacatttcac actcagctct ga 22

<210> 124
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 124
 caggagcatt tcgtgaaaga 20

<210> 125
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 125
 ttttgacct ttatctgcag cctttg 26

<210> 126
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 126
 tttaaccga gcttcctcat 20

<210> 127
 <211> 22

<212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 127
 ctgcaaaatc ttacgacttt gg

22

<210> 128
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 128
 caacaacaa tggctacatc aaatttagca

30

<210> 129
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 129
 atgacactca gcaaacctga gt

22

<210> 130
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 130
 ctgcaaaatc ttacgacttt gg

22

<210> 131
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer
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<400> 131
 caacaacaa tggctacatc aaatttagca

30

<210> 132
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 132
 atgacactca gcaaacctga gt 22

<210> 133
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 133
 ctgcaaaatc ttacgacttt gg 22

<210> 134
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 134
 caacaacaa tggctacatc aaatttagca 30

<210> 135
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 135
 tcagcaaacc tgagtcctgt a 21

<210> 136
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 136
 ctgcaaaatc ttacgacttt gg 22

<210> 137
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

 <400> 137
 caacaaacaa tggctacatc aaatttagca 30

<210> 138
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

 <400> 138
 atgacactca gcaaacctga gt 22

<210> 139
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

 <400> 139
 ctgcaaaatc ttacgacttt gg 22

<210> 140
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

 <400> 140
 caacaaacaa tggctacatc aaatttagca 30

<210> 141
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 141
atgacactca gcaaacctga gt 22

<210> 142
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 142
gggctataag tcagtcggaa gt 22

<210> 143
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 143
cctgtatttg tctgccaagc caatcg 26

<210> 144
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 144
acagtcgaga ggaacacaca tc 22

<210> 145
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

<400> 145
gaggacagct ttgatttcac tg 22

<210> 146
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR Primer
sequence

sequence

<400> 146
 tggatttgat ccatttcctc tctacca 27

<210> 147
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PCR Primer
 sequence

<400> 147
 aagagactgg atggcttttc at 22

<210> 148
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
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<400> 149
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<210> 150
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<210> 153
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<210> 154
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<210> 157

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<210> 159

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<210> 169
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<210> 172
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<210> 173
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<210> 176
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<400> 176
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<210> 177
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<210> 178
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<220>
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<400> 178
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<210> 179
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<220>
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<400> 179
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<210> 180
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<220>

<223> Description of Artificial Sequence: PCR Primer
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<400> 180
ttatgatgtc ccagagcttg tc 22

<210> 181
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<400> 181
gttctgtgtg gtcataatc ct 22

<210> 182
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<220>
<223> Description of Artificial Sequence: PCR Primer
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<210> 183
<211> 20
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<220>
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<400> 183
cttgcccttg tacatttcca 20

<210> 184
<211> 22
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<400> 184
caattgcctc cagtatttga ac 22

<210> 185
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<220>
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<400> 185
 ttgcagacat agggtaacct cacatt 26

<210> 186
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<220>
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<400> 186
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<210> 187
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<220>
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<400> 187
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<210> 188
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<400> 188
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<210> 189
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<400> 189
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<210> 190
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 <400> 190
 ggaggccaca ggagcaggat ca 22

 <210> 191
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 <220>
 <223> Description of Artificial Sequence: PCR Primer
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 <400> 191
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 <210> 192
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 <400> 192
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 <210> 193
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 <400> 193
 ggatccagcc ctggccaggc cgtgtgcaac ttcg 34

 <210> 194
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 <400> 194
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<210> 195
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 <400> 195
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<210> 196
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 <400> 196
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<210> 197
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 <400> 197
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<210> 198
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 <400> 198
 ccaaggttga ccacctccat 20

<210> 199
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 <400> 199
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<210> 200
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<400> 200
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<210> 201
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 <223> Description of Artificial Sequence: PCR Primer
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<400> 201
 ccgcctgtgt tccatggctt 20

<210> 202
 <211> 23
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<220>
 <223> Description of Artificial Sequence: PCR Primer
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<400> 202
 gtcattctgc tgccggttg tag 23

<210> 203
 <211> 24
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<220>
 <223> Description of Artificial Sequence: PCR Primer
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<400> 203
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<210> 204
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<220>
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<400> 204
ttacaattgc ctccagtatt tgaacttgca 30

<210> 205
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<400> 205
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<400> 206
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<400> 207
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<223> Description of Artificial Sequence: PCR Primer
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<400> 208
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sequence

<400> 209
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 aaaaaaaaaa aaaaaaaaaa 3260

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 Pro Ile Val Ile Ser Val Ser Gln Ser Thr Val Tyr Ile Asn Asn Lys
 35 40 45
 Ala Lys Ile Ile Ser Ser Asp Ile Ile Ser Thr Asn Gly Ile Val His
 50 55 60
 Ile Ile Asp Lys Leu Leu Ser Pro Lys Asn Leu Leu Ile Thr Pro Lys
 65 70 75 80
 Asp Asn Ser Gly Arg Ile Leu Gln Asn Leu Thr Thr Leu Ala Thr Asn
 85 90 95
 Asn Gly Tyr Ile Lys Phe Ser Asn Leu Ile Gln Asp Ser Gly Leu Leu
 100 105 110
 Ser Val Ile Thr Asp Pro Ile His Thr Pro Val Thr Leu Phe Trp Pro
 115 120 125
 Thr Asp Gln Ala Leu His Ala Leu Pro Ala Glu Gln Gln Asp Phe Leu
 130 135 140
 Phe Asn Gln Asp Asn Lys Asp Lys Leu Lys Glu Tyr Leu Lys Phe His
 145 150 155 160
 Val Ile Arg Asp Ala Lys Val Leu Ala Val Asp Leu Pro Thr Ser Thr
 165 170 175
 Ala Trp Lys Thr Leu Gln Gly Ser Glu Leu Ser Val Lys Cys Gly Ala
 180 185 190
 Gly Arg Asp Ile Gly Asp Leu Phe Leu Asn Gly Gln Thr Cys Arg Ile
 195 200 205
 Val Gln Arg Glu Leu Leu Phe Asp Leu Gly Val Ala Tyr Gly Ile Asp
 210 215 220
 Cys Leu Leu Ile Asp Pro Thr Leu Gly Gly Arg Cys Asp Thr Phe Thr
 225 230 235 240
 Thr Phe Asp Ala Ser Gly Glu Cys Gly Ser Cys Val Asn Thr Pro Ser
 245 250 255
 Cys Pro Arg Trp Ser Lys Pro Lys Gly Val Lys Gln Lys Cys Leu Tyr
 260 265 270

Asn Leu Pro Phe Lys Arg Asn Leu Glu Gly Cys Arg Glu Arg Cys Ser
 275 280 285
 Leu Val Ile Gln Ile Pro Arg Cys Cys Lys Gly Tyr Phe Gly Arg Asp
 290 295 300
 Cys Gln Ala Cys Pro Gly Gly Pro Asp Ala Pro Cys Asn Asn Arg Gly
 305 310 315 320
 Val Cys Leu Asp Gln Tyr Ser Ala Thr Gly Glu Cys Lys Cys Asn Thr
 325 330 335
 Gly Phe Asn Gly Thr Ala Cys Glu Met Cys Trp Pro Gly Arg Phe Gly
 340 345 350
 Pro Asp Cys Leu Pro Cys Gly Cys Ser Asp His Gly Gln Cys Asp Asp
 355 360 365
 Gly Ile Thr Gly Ser Gly Gln Cys Leu Cys Glu Thr Gly Trp Thr Gly
 370 375 380
 Pro Ser Cys Asp Thr Gln Ala Val Leu Pro Ala Val Cys Thr Pro Pro
 385 390 395 400
 Cys Ser Ala His Ala Thr Cys Lys Glu Asn Asn Thr Cys Glu Cys Asn
 405 410 415
 Leu Asp Tyr Glu Gly Asp Gly Ile Thr Cys Thr Val Val Asp Phe Cys
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 435 440 445
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 450 455 460
 His Ser Cys Thr Glu Ile Asp Pro Cys Ala Asp Gly Leu Asn Gly Gly
 465 470 475 480
 Cys His Glu His Ala Thr Cys Lys Met Thr Gly Pro Gly Lys His Lys
 485 490 495
 Cys Glu Cys Lys Ser His Tyr Val Gly Asp Gly Leu Asn Cys Glu Pro
 500 505 510
 Glu Gln Leu Pro Ile Asp Arg Cys Leu Gln Asp Asn Gly Gln Cys His
 515 520 525
 Ala Asp Ala Lys Cys Val Asp Leu His Phe Gln Asp Thr Thr Val Gly
 530 535 540
 Val Phe His Leu Arg Ser Pro Leu Gly Gln Tyr Lys Leu Thr Phe Asp
 545 550 555 560
 Lys Ala Arg Glu Ala Cys Ala Asn Glu Ala Ala Thr Met Ala Thr Tyr
 565 570 575
 Asn Gln Leu Ser Tyr Ala Gln Lys Ala Lys Tyr His Leu Cys Ser Ala
 580 585 590
 Gly Trp Leu Glu Thr Gly Arg Val Ala Tyr Pro Thr Ala Phe Ala Ser
 595 600 605

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Gln Asn Cys Gly Ser Gly Val Val Gly Ile Val Asp Tyr Gly Pro Arg
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 Pro Asn Lys Ser Glu Met Trp Asp Val Phe Cys Tyr Arg Met Lys Asp
 625 630 635 640
 Val Asn Cys Thr Cys Lys Val Gly Tyr Val Gly Asp Gly Phe Ser Cys
 645 650 655
 Ser Gly Asn Leu Leu Gln Val Leu Met Ser Phe Pro Ser Leu Thr Asn
 660 665 670
 Phe Leu Thr Glu Val Leu Ala Tyr Ser Asn Ser Ser Ala Arg Gly Arg
 675 680 685
 Ala Phe Leu Glu His Leu Thr Asp Leu Ser Ile Arg Gly Thr Leu Phe
 690 695 700
 Val Pro Gln Asn Ser Gly Leu Gly Glu Asn Glu Thr Leu Ser Gly Arg
 705 710 715 720
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 725 730 735
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 740 745 750
 Ile Thr Ala Ser Gln Asp Pro Leu Gln Pro Thr Glu Thr Arg Phe Val
 755 760 765
 Asp Gly Arg Ala Ile Leu Gln Trp Asp Ile Phe Ala Ser Asn Gly Ile
 770 775 780
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 785 790 795 800
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 820 825 830
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 835 840 845
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 865 870 875 880
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